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(54) Title: COMBINATION METHODS OF INHIBITING TUMOR GROWTH WITH A VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPTOR ANTAGONIST

(57) Abstract: The present invention provides a method of reducing or inhibiting tumor growth in a mammal comprising treating the mammal with an effective amount of a combination of a VEGF receptor antagonist and radiation, chemotherapy, and/or an additional receptor antagonist.



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**COMBINATION METHODS OF INHIBITING TUMOR GROWTH  
WITH A VASCULAR ENDOTHELIAL GROWTH FACTOR  
RECEPTOR ANTAGONIST**

- [01] This application is a continuation-in-part of Application No. 09/798,689, filed March 2, 2001, pending, which is a continuation-in-part of Application No. 09/401,163, filed on September 22, 1999, pending, which is a continuation of Application No. 08/967,113 filed on November 10, 1997, pending, which is a continuation-in-part of Patent Number 5,861,499 filed September 3, 1996, which is a continuation-in-part of Application No. 08/476,533 filed June 7, 1995, abandoned, which is a continuation of Patent Number 5,840,301 filed October 20, 1994, which is a continuation-in-part of Application No. 08/196,041 filed February 10, 1994, abandoned. The entire disclosures of the aforementioned prior applications are incorporated herein by reference.

**FIELD OF THE INVENTION**

- [02] The present invention is directed to methods of treating tumors utilizing vascular endothelial growth factor (VEGF) receptor antagonists in combination with a chemotherapeutic agent, radiation, and/or a different growth factor receptor antagonist.

**BACKGROUND OF THE INVENTION**

- [03] Angiogenesis, which refers to the formation of capillaries from pre-existing vessels in the embryo and adult organism, is known to be a key element in tumor growth, survival and metastasis. Growth factors and their receptors, including epidermal growth factor (EGF), transforming growth factor- $\alpha$  (TGF- $\alpha$ ), transforming growth factor- $\beta$  (TGF- $\beta$ ), acidic and basic fibroblast growth factor (aFGF and bFGF), platelet derived growth factor (PDGF), and vascular endothelial growth factor (VEGF), are thought to play a role in tumor angiogenesis. *See Klagsbrun & D'Amore, Annual Rev. Physiol.*, 53: 217-239 (1991). Binding of these growth factors to their cell surface receptors induces receptor activation, which initiates and modifies signal transduction pathways and leads to cell proliferation and differentiation. VEGF, an endothelial cell-specific mitogen, is distinct among these factors in that it acts as an angiogenesis inducer by specifically promoting the proliferation of endothelial cells.

- [04] VEGF is a homodimeric glycoprotein consisting of two 23 kD subunits and is a strong inducer of vascular permeability, stimulator of endothelial cell migration and proliferation, and an important survival factor for newly formed blood vessels. Four different monomeric isoforms of VEGF exist, resulting from alternative splicing of mRNA. These include two membrane bound forms (VEGF<sub>206</sub> and VEGF<sub>189</sub>) and two soluble forms (VEGF<sub>165</sub> and VEGF<sub>121</sub>). In all human tissues except placenta, VEGF<sub>165</sub> is the most abundant isoform.
- [05] VEGF is a key regulator of vasculogenesis, which is the *de novo* development of new blood vessels from the differentiation of endothelial precursors (angioblasts) *in situ*, and is expressed in embryonic tissues (Breier et al., Development (Camb.), 114:521 (1992)), macrophages, proliferating epidermal keratinocytes during wound healing (Brown et al., J. Exp. Med., 176:1375 (1992)), and may be responsible for tissue edema associated with inflammation (Ferrara et al., Endocr. Rev., 13:18 (1992)). *In situ* hybridization studies have demonstrated high VEGF expression in a number of human tumor lines including glioblastoma multiforme, hemangioblastoma, central nervous system neoplasms and AIDS-associated Kaposi's sarcoma (Plate et al. (1992) Nature 359: 845-848; Plate et al. (1993) Cancer Res. 53: 5822-5827; Berkman et al. (1993) J. Clin. Invest. 91: 153-159; Nakamura et al. (1992) AIDS Weekly, 13 (1)). High levels of VEGF were also observed in hypoxia induced angiogenesis (Shweiki et al. (1992) Nature 359: 843-845).
- [06] The biological response of VEGF is mediated through its high affinity receptors, which are selectively expressed on endothelial cells during embryogenesis (Millauer, Cell, 72: 835-846 (1993)) and during tumor formation. VEGF receptors (VEGFRs) typically are class III receptor-type tyrosine kinases characterized by having several, typically 5 or 7, immunoglobulin-like loops in their amino-terminal extracellular receptor ligand-binding domains (Kaipainen et al., J. Exp. Med., 178:2077-2088 (1993)). The other two regions include a transmembrane region and a carboxy-terminal intracellular catalytic domain interrupted by an insertion of hydrophilic interkinase sequences of variable lengths, called the kinase insert domain (Terman et al., Oncogene, 6:1677-1683 (1991)). VEGFRs include *fms*-like tyrosine kinase receptor (flt-1), or VEGFR-1, sequenced by Shibuya et al., Oncogene, 5: 519-524 (1990), kinase insert domain-containing receptor/fetal liver kinase (KDR/flk-1), or VEGFR-2, described in WO 92/14248, filed February 20, 1992, and

Terman et al., *Oncogene*, 6: 1677-1683 (1991) and sequenced by Matthews et al., *Proc. Natl. Acad. Sci. USA*, 88: 9026-9030 (1991), although other receptors, such as neuropilin-1 and -2, can also bind VEGF. Another tyrosine kinase receptor, VEGFR-3 (flt-4), binds the VEGF homologues VEGF-C and VEGF-D and is more important in the development of lymphatic vessels.

[07] High levels of Flk-1 are expressed by endothelial cells that infiltrate gliomas (Plate et al., (1992) *Nature* 359: 845-848). Flk-1 levels are specifically upregulated by VEGF produced by human glioblastomas (Plate et al. (1993) *Cancer Res.* 53: 5822-5827). The finding of high levels of Flk-1 expression in glioblastoma associated endothelial cells (GAEC) indicates that receptor activity is probably induced during tumor formation since Flk-1 transcripts are barely detectable in normal brain endothelial cells. This upregulation is confined to the vascular endothelial cells in close proximity to the tumor. Blocking VEGF activity with neutralizing anti-VEGF monoclonal antibodies (mAbs) resulted in an inhibition of the growth of human tumor xenografts in nude mice (Kim et al. (1993) *Nature* 362: 841-844), indicating a direct role for VEGF in tumor-related angiogenesis.

[08] Although the VEGF ligand is upregulated in tumor cells, and its receptors are upregulated in tumor infiltrated vascular endothelial cells, the expression of the VEGF ligand and its receptors is low in normal cells that are not associated with angiogenesis. Therefore, such normal cells would not be affected by blocking the interaction between VEGF and its receptors to inhibit angiogenesis, and therefore tumor growth.

[09] An object of the present invention is to provide VEGF receptor antagonists. A further object of this invention is to provide methods to inhibit angiogenesis and thereby to inhibit or reduce tumor growth in mammals using such VEGF receptor antagonists and, in particular, using such VEGF receptor antagonists combined with radiation, chemotherapy, or another receptor antagonist.

#### BRIEF SUMMARY OF THE INVENTION

[10] The present invention provides methods of reducing or inhibiting tumor growth in a mammal by administering an effective amount of a combination of a VEGF receptor antagonist and another receptor antagonist. Also provided by the present invention are methods of reducing or inhibiting tumor growth in a mammal by administering an

effective amount of a combination of a VEGF receptor antagonist and radiation. In addition, the present invention provides methods of reducing or inhibiting tumor growth in a mammal by administering an effective amount of a combination of a VEGF receptor antagonist and a chemotherapeutic agent.

#### BRIEF DESCRIPTION OF THE FIGURES

- [11] **Figure 1:** Western Blot of flk-1 (VEGFR-2)/SEAPS immunoprecipitation with monoclonal antibody DC-101 demonstrating that DC-101 immunoprecipitates murine flk-1:SEAPS but not SEAPS alone.
- [12] **Figures 2a and 2b:** **Figure 2a:** Competitive inhibition assay indicating the effect of anti-flk-1 (VEGFR-2) monoclonal antibody DC-101 on VEGF<sub>165</sub> induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor in transfected 3T3 cells. **Figure 2b:** Sensitivity of VEGF induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor to inhibition by monoclonal antibody DC-101. C441 cells were assayed at maximal stimulatory concentrations of VEGF<sub>165</sub> (40 ng/ml) combined with varying levels of the antibody.
- [13] **Figures 3a and 3b:** **Figure 3a:** Titration of VEGF-induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor in the presence of mAb DC-101. C441 cells were stimulated with the concentrations of VEGF indicated in the presence (Lanes 1 to 4) or absence (Lanes 5 to 8) of 5 µg/ml of mAb DC-101. Unstimulated cells assayed in the presence of antibody (Lane 9) serves as the control. **Figure 3b:** Densitometry scans of the level of phosphorylated receptor in each lane in Figure 3a relative to each VEGF concentration is plotted to show the extent of mAb inhibition at excess ligand concentrations. Cell lysates were prepared for detection by anti-phosphotyrosine as described in the Examples below.
- [14] **Figure 4:** Inhibition of VEGF-flk-1 (VEGFR-2)/fms activation by prebound mAb DC-101. C441 cells were stimulated with the concentrations of VEGF indicated in the absence (Lanes 3 and 4) and presence (Lanes 5 and 6) of DC-101. Unstimulated cells (Lanes 1 and 2) serve as controls. MAb was assayed using two sets of conditions. For P, cells were prebound with mAb followed by stimulation with VEGF for 15 minutes at

room temperature. For C, mAb and ligand were added simultaneously and assayed as above.

- [15] **Figure 5:** VEGF-induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor by treatments with varying concentrations of monoclonal antibody DC-101 and conditioned media from glioblastoma cells (GB CM).
- [16] **Figure 6:** FACS analysis of anti-flk-1 (VEGFR-2) mAb binding to flk-1 (VEGFR-2)/fms transfected 3T3 Cells (C441). Transfected flk-1 (VEGFR-2)/fms 3T3 cells were incubated on ice for 60 minutes with 10 µg/ml of the anti-flk-1 (VEGFR-2) mAb DC-101 or the isotype matched irrelevant anti-flk-1 mAb 23H7. Cells were washed and reincubated with 5 µg of goat anti-mouse IgG conjugated to FITC, washed, and analyzed by flow cytometry to determine antibody binding. Data shows the level of fluorescence for DC-101 to C441 cells relative to that detected with the irrelevant mAb 23H7.
- [17] **Figure 7:** Saturation binding of mAb DC-101 to the flk-1 (VEGFR-2)/fms receptor on the transfected 3T3 cell line C441. Confluent C441 cells were incubated in 24 well plates with increasing concentrations of mAb DC-101 (50 ng/ml to 2 µg/ml) for two hours at 4°C. Cells were washed and incubated with 5 µg anti-rat IgG-biotin conjugate. To detect binding, cells were washed, incubated with a 1:1000 dilution of streptavidin-HRP, washed and incubated in a colormetric detection system (TMB). Data represents the absorbance at 540 nm versus increasing concentrations of mAb DC-101. The binding of the secondary antibody to cells alone was subtracted from each determination to adjust for non-specific binding. Data represents the average of three independent experiments.
- [18] **Figure 8:** Immunoprecipitation of phosphorylated flk-1 (VEGFR-2)/fms from VEGF stimulated flk-1 (VEGFR-2)/fms transfected 3T3 cells. Cells were stimulated with VEGF as described in the Experimental Procedures and lysates were immunoprecipitated with irrelevant or relevant antibodies as follows: 1. rat anti-*FLK2* IgG2a (mAb 2A13); 2. rat anti-flk-1 (VEGFR-2) IgG1 (mAb DC-101); 3. rat anti-*FLK2* IgG1 (mAb 23H7); 4. rabbit anti-fms polyclonal antibody. Immunoprecipitated protein was subjected to SDS PAGE followed by Western blotting. The immunoprecipitation of VEGF activated receptor was detected by probing the blots with an anti-phosphotyrosine antibody.

- [19] **Figure 9:** Sensitivity of VEGF-induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor to inhibition by mAb DC-101. Prebound and competitive assays were performed with 40 ng/ml of VEGF at the antibody concentrations indicated. Cell lysates were prepared for receptor detection with anti-phosphotyrosine as described in the Examples below.
- [20] **Figure 10:** Effect of mAb DC-101 on CSF-1 induced phosphorylation of the fms receptor. In (B), the fms/*FLK-2* transfected 3T3 cell line, 10A2, was stimulated with optimal stimulatory levels of CSF-1 in the absence (Lanes 3 and 4) and presence (Lanes 5 and 6) of 5 µg/ml of mAb DC-101. Unstimulated cells assayed in the absence (Lane 1) or presence (Lane 2) of antibody serve as controls. Cell lysates were prepared for detection by anti-phosphotyrosine as described in the Examples below.
- [21] **Figure 11:** Specificity of mAb DC-101 neutralization of the activated flk-1 (VEGFR-2)/fms receptor. C441 cells were stimulated with 20 or 40 ng/ml of VEGF in the presence of DC-101 (IgG1) or the irrelevant anti-*FLK-2* rat monoclonal antibodies 2A13 (IgG2a) or 23H7 (IgG1). Assays were performed with each antibody in the absence of VEGF (Lanes 1 to 3) and in the presence of VEGF under competitive (lanes 4 to 8) or prebound (lanes 9 to 11) conditions. Cell lysates were prepared for detection by anti-phosphotyrosine as described in the Examples below. Blots were stripped and reprobed to detect the flk-1 (VEGFR-2)/fms receptor using a rabbit polyclonal antibody to the C-terminal region of the fms receptor.
- [22] **Figure 12:** Immunoprecipitation of phosphorylated receptor bands from VEGF stimulated HUVEC cells. HUVEC cells were grown to subconfluency in endothelial growth medium (EGM) for three days without a change of medium. Receptor forms were immunoprecipitated by mAb DC-101 from lysates of unstimulated cells (Lane 1), VEGF stimulated cells (lane 2), and cells stimulated with VEGF in the presence of 1 µg/ml heparin (Lane 3). Phosphorylation assays, immunoprecipitations, and detection of the phosphorylated receptor forms were performed as described in the Experimental Procedures.
- [23] **Figure 13:** Effect of mAb DC-101 on the proliferation of HUVEC cells in response to VEGF. Cells were grown for 48 hours as described in the legend to Figure 6. Cells were then subjected to the following assay conditions: no addition to medium

(untreated); a change of fresh endothelial growth medium (complete medium); the addition of 10 ng/ml of VEGF in the absence or presence of 1  $\mu$ g/ml heparin; and VEGF and VEGF-heparin treated cells assayed in the presence of 1  $\mu$ g/ml of DC-101. Cells were assayed for proliferation by colormetric detection at 550 nm using a cell proliferation assay kit (Promega).

- [24]       **Figures 14a and 14b** **Figure 14a:** Reduction in tumor growth of individual animals with DC-101 (rat anti-flk-1 monoclonal antibody). **Figure 14b:** Reduction in tumor growth in individual animals with the control 2A13 group (rat anti-flk-2 monoclonal antibody).
- [25]       **Figure 15:** Athymic nude mice were injected subcutaneously with human glioblastoma cell line GBM-18 and divided into three groups: a PBS control, an irrelevant rat IgG1 control 23H7, and DC-101. Treatments were administered simultaneously with tumor xenografts and continued for four weeks.
- [26]       **Figure 16:** A graph showing the direct binding of different scFv antibodies (p1C11, p1F12, p2A6 and p2A7) to immobilized KDR (VEGFR-2).
- [27]       **Figure 17:** A graph showing the inhibition of binding of KDR (VEGFR-2) to immobilized VEGF<sub>165</sub> by different scFv antibodies (p1C11, p1F12, p2A6 and p2A7).
- [28]       **Figure 18:** A graph showing the inhibition of VEGF-induced HUVEC proliferation by scFv antibodies (p2A6 and p1C11).
- [29]       **Figure 19:** The nucleotide and deduced amino acid sequence of V<sub>H</sub> and V<sub>L</sub> chains of c-p1C11.
- [30]       **Figure 20:** A graph showing the direct binding of antibodies (c-p1C11, p1C11, p2A6) to immobilized KDR (VEGFR-2).
- [31]       **Figure 21:** A graph showing the FACS analysis of c-p1C11 binding to KDR- (VEGFR-2) expressing HUVEC.
- [32]       **Figure 22:** A graph showing the inhibition of binding of KDR (VEGFR-2) receptor to immobilized VEGF<sub>165</sub> by different scFv antibodies (c-p1C11, p1C11, and p2A6).

[33] **Figure 23:** A graph showing the inhibition of binding of radiolabeled VEGF<sub>165</sub> to immobilized KDR (VEGFR-2) receptor by c-p1C11 and cold VEGF<sub>165</sub>.

[34] **Figure 24:** A graph showing the inhibition of VEGF-induced HUVEC proliferation by anti-KDR (VEGFR-2) antibodies (c-p1C11, p1C11).

#### DETAILED DESCRIPTION OF THE INVENTION

[35] The present invention provides methods of reducing or inhibiting tumor growth in mammals with radiation, chemotherapy, and/or an additional receptor antagonist in combination with VEGF receptor antagonists.

[36] In a preferred embodiment, there is synergy when tumors, including human tumors, are treated with a VEGF receptor antagonist in conjunction with chemotherapeutic agents, radiation, or an additional receptor antagonist or combinations thereof. In other words, the inhibition of tumor growth by a VEGF receptor antagonist is enhanced more than expected when combined with chemotherapeutic agents, radiation, or an additional receptor antagonist or combinations thereof. Synergy may be shown, for example, by greater inhibition of tumor growth with combined treatment than would be expected from the additive effect of treatment with a VEGF receptor antagonist and a chemotherapeutic agent, radiation, or an additional receptor antagonist. Preferably, synergy is demonstrated by remission of the cancer where remission is not expected from treatment with a combination of a VEGF receptor antagonist and a chemotherapeutic agent, radiation, or an additional receptor antagonist. (See Example VIII.)

[37] The VEGF receptor antagonist is administered before, during, or after commencing chemotherapy or radiation therapy, as well as any combination thereof, i.e. before and during, before and after, during and after, or before, during, and after commencing the chemotherapy and/or radiation therapy. For example, when the VEGF receptor antagonist is an antibody, the antibody is typically administered between 1 and 30 days, preferably between 3 and 20 days, more preferably between 5 and 12 days before commencing radiation therapy and/or chemotherapy.

[38] *Radiation*

[39] The source of radiation, used in combination with a VEGF receptor antagonist, can be either external or internal to the patient being treated. When the source is external to the patient, the therapy is known as external beam radiation therapy (EBRT). When the source of radiation is internal to the patient, the treatment is called brachytherapy (BT).

[40] The radiation is administered in accordance with well known standard techniques using standard equipment manufactured for this purpose, such as AECL Theratron and Varian Clinac. The dose of radiation depends on numerous factors as is well known in the art. Such factors include the organ being treated, the healthy organs in the path of the radiation that might inadvertently be adversely affected, the tolerance of the patient for radiation therapy, and the area of the body in need of treatment. The dose will typically be between 1 and 100 Gy, and more particularly between 2 and 80 Gy. Some doses that have been reported include 35 Gy to the spinal cord, 15 Gy to the kidneys, 20 Gy to the liver, and 65-80 Gy to the prostate. It should be emphasized, however, that the invention is not limited to any particular dose. The dose will be determined by the treating physician in accordance with the particular factors in a given situation, including the factors mentioned above.

[41] The distance between the source of the external radiation and the point of entry into the patient may be any distance that represents an acceptable balance between killing target cells and minimizing side effects. Typically, the source of the external radiation is between 70 and 100 cm from the point of entry into the patient.

[42] Brachytherapy is generally carried out by placing the source of radiation in the patient. Typically, the source of radiation is placed approximately 0-3 cm from the tissue being treated. Known techniques include interstitial, intercavitary, and surface brachytherapy. The radioactive seeds can be implanted permanently or temporarily. Some typical radioactive atoms that have been used in permanent implants include iodine-125 and radon. Some typical radioactive atoms that have been used in temporary implants include radium, cesium-137, and iridium-192. Some additional radioactive atoms that have been used in brachytherapy include americium-241 and gold-198.

- [43] The dose of radiation for brachytherapy can be the same as that mentioned above for external beam radiation therapy. In addition to the factors mentioned above for determining the dose of external beam radiation therapy, the nature of the radioactive atom used is also taken into account in determining the dose of brachytherapy.
- [44] *Chemotherapy*
- [45] Chemotherapeutic agents include all chemical compounds that are effective in inhibiting tumor growth.
- [46] The administration of chemotherapeutic agents can be accomplished in a variety of ways including systemically by the parenteral and enteral routes. In one embodiment, the VEGF receptor antagonist and the chemotherapeutic agent are administered as separate molecules. In another embodiment, the VEGF receptor antagonist is attached, such as, for example, by conjugation, to a chemotherapeutic agent.
- [47] Examples of chemotherapeutic agents include alkylating agents, for example, nitrogen mustards, ethyleneimine compounds and alkyl sulphonates; antimetabolites, for example, folic acid, purine or pyrimidine antagonists; mitotic inhibitors, for example, vinca alkaloids and derivatives of podophyllotoxin; cytotoxic antibiotics; compounds that damage or interfere with DNA expression.
- [48] Additionally, chemotherapeutic agents include antibodies, biological molecules and small molecules, as described herein.
- [49] Particular examples of chemotherapeutic agents or chemotherapy include cisplatin, dacarbazine (DTIC), dactinomycin, mechlorethamine (nitrogen mustard), streptozocin, cyclophosphamide, carmustine (BCNU), lomustine (CCNU), doxorubicin (adriamycin), daunorubicin, procarbazine, mitomycin, cytarabine, etoposide, methotrexate, 5-fluorouracil, vinblastine, vincristine, bleomycin, paclitaxel (taxol), docetaxel (taxotere), aldesleukin, asparaginase, busulfan, carboplatin, cladribine, dacarbazine, floxuridine, fludarabine, hydroxyurea, ifosfamide, interferon alpha, leuprolide, megestrol, melphalan, mercaptopurine, plicamycin, mitotane, pegaspargase, pentostatin, pipobroman, plicamycin, streptozocin, tamoxifen, teniposide, testolactone, thioguanine, thiotepa, uracil mustard, vinorelbine, chlorambucil, taxol and combinations thereof.

**[50]**    *Growth Factor Receptor Antagonists*

**[51]**            Growth factor receptor antagonists (other than VEGF receptor antagonists) that can be used in the context of the present invention include all substances that inhibit the stimulation of a growth factor receptor by a growth factor receptor ligand. Such inhibition of stimulation inhibits the growth of cells that express the growth factor receptor.

**[52]**            Some examples of growth factor receptors involved in tumorigenesis are the receptors for epidermal growth factor (EGFR), platelet-derived growth factor (PDGFR), insulin-like growth factor (IGFR), nerve growth factor (NGFR), and fibroblast growth factor (FGF).

**[53]**            Preferably, the growth factor receptor antagonist to be used in this invention is an EGFR antagonist. An EGFR antagonist, in the context of the present invention, is a biological molecule, small molecule, or any other substance that inhibits EGFR activation, thereby inhibiting the tyrosine kinase activity of EGFR, and preventing receptor autophosphorylation and the phosphorylation of other proteins involved in the various EGFR signaling pathways. By inhibition of activation of EGFR is meant any decrease in the activation of the EGFR, which need not completely prevent or stop activation of EGFR.

**[54]**            Moreover, inhibition of EGFR activation, as defined by the present invention, means inhibition of EGFR resulting from interaction of the EGFR antagonist and the receptor. By interaction is meant sufficient physical or chemical interaction between the EGFR antagonist and the receptor, such that tyrosine kinase activity is inhibited. One of skill in the art would appreciate that examples of such chemical interactions, which include association or bonding, are known in the art and include covalent bonding, ionic bonding, hydrogen bonding, etc., between the EGFR antagonist and the receptor. This is in contrast with an EGF antagonist, which interacts with the ligand, thereby inhibiting activation.

**[55]**            As is the case with other growth factors, increased EGFR activation can result from higher levels of ligand, EGFR gene amplification, increased transcription of the receptor or mutations that cause unregulated receptor signaling. Amplification of the gene encoding EGFR results in an increased number of ligands binding to the EGFR, which can

further stimulate cell proliferation. EGFR may also be overexpressed in the absence of gene amplification, presumably through mutations that increase EGFR transcription, mRNA translation, or stability of the protein. EGFR mutants have been identified in gliomas, non-small-cell lung carcinomas, ovarian carcinomas and prostate carcinomas that have a constitutively active tyrosine kinase, suggesting a role for high-level EGFR activity rather than EGFR overexpression in these cancers. *See, e.g., Pedersen et al., Ann. Oncol., 12(6):745-60 (2001).* (Type III EGFR mutation – variously named EGFRvIII, de2-7 EGFR or AEGFR – lacks a portion of the extracellular ligand binding domain encoded by exons 2-7.); *see also Wikstrand et al., Cancer Res., 55:3140-3148 (1995).*

[56] In one embodiment of the present invention, the EGFR antagonist inhibits binding of EGFR to its ligand. Binding of a ligand to an external, extracellular domain of EGFR stimulates receptor dimerization, autophosphorylation of EGFR, activation of the receptor's internal, cytoplasmic tyrosine kinase domain, and initiation of multiple signal transduction pathways involved in regulation of DNA synthesis and cell division. Ligands for EGFR include, for example, EGF, TGF- $\alpha$ , amphiregulin, heparin-binding EGF (HB-EGF) and betarecullulin. EGF and TGF- $\alpha$  are thought to be the main endogenous ligands that result in EGFR-mediated stimulation, although TGF- $\alpha$  has been shown to be more potent in promoting angiogenesis.

[57] In another embodiment of the present invention, the EGFR antagonist binds EGFR. It should be appreciated that the EGFR antagonist can bind externally to the extracellular portion of EGFR, which may or may not inhibit binding of the ligand, or internally to the tyrosine kinase domain. Examples of EGFR antagonists that bind EGFR include, without limitation, biological molecules, such as antibodies (and functional equivalents thereof) specific for EGFR, and synthetic kinase inhibitors that act directly on the cytoplasmic domain of EGFR, such as small molecules.

[58] The EGFR antagonist of the present invention is preferably a biological molecule, more preferably an antibody, or functional equivalent thereof, specific for EGFR. A description of the antibodies useful in the present invention can be found in the section entitled "Antibodies." Furthermore, following antibody binding, the EGFR-antibody complex is preferably internalized and degraded, preventing receptor re-utilization by the cell.

- [59] A known biological molecule EGFR antagonist is ERBITUX<sup>TM</sup> (IMC-C225), which is a chimeric (human/mouse) monoclonal antibody specific for EGFR. *See, e.g.*, U.S. Patent No. 4,943,533 (Mendelsohn et al.); U.S. Patent No. 6,217,866 (Schlessinger et al.); U.S. Application Nos. 08/973,065 (Goldstein et al.) and 09/635,974 (Teufel); WO 99/60023 (Waksal et al.) and WO 00/69459 (Waksal). The monoclonal antibody ERBITUX<sup>TM</sup> specifically binds EGFR and blocks binding of a ligand, e.g., EGF. This blockade results in inhibition of tumor growth, which includes inhibition of tumor invasion, metastases, cell repair and angiogenesis, by interfering with the effects of EGFR activation. In addition, or alternatively, the monoclonal antibody ERBITUX<sup>TM</sup> may promote internalization of the receptor-antibody complex, preventing further stimulation of the receptor by its ligand or any other mechanism.
- [60] Another example of a biological molecule EGFR antagonist is ABX-EGF, which is a fully human IgG<sub>2</sub> monoclonal antibody specific for EGFR. ABX-EGF binds EGFR with high specificity, blocking binding of EGFR to both of its ligands, EGF and TGF- $\alpha$ . *See, e.g.*, Figlin et al., Abstract 1102 presented at the 37th Annual Meeting of ASCO, San Francisco, CA, 12-15 May 2001. The sequence and characterization of ABX-EGF, which was formerly known as clone E7.6.3, is disclosed in U.S. Patent No. 6,235,883 (Abgenix, Inc.) at col. 28, line 62 through col. 29, line 36 and in Fig. 29-34. *See* Yang et al., *Critical Rev. Oncol./Hematol.*, 38(1): 17-23, 2001.
- [61] In an alternative, but also preferred, embodiment, the EGFR antagonist of the present invention is a small molecule tyrosine kinase inhibitor. A description of small molecules can be found in the section entitled "Small Molecules." Numerous small molecules have been described as being useful to inhibit EGFR.
- [62] For example, Spada et al., U.S. Patent 5,656,655, discloses styryl substituted heteroaryl compounds that inhibit EGFR. The heteroaryl group is a monocyclic ring with one or two heteroatoms, or a bicyclic ring with 1 to about 4 heteroatoms, the compound being optionally substituted or polysubstituted. The compounds disclosed in U.S. Patent 5,656,655 are incorporated herein by reference.

- [63] Spada et al., U.S. Patent 5,646,153 discloses bis mono and/or bicyclic aryl heteroaryl, carbocyclic, and heterocarbocyclic compounds that inhibit EGFR. The compounds disclosed in U.S. Patent 5,646,153 are incorporated herein by reference.
- [64] Bridges et al., U.S. Patent 5,679,683 discloses tricyclic pyrimidine compounds that inhibit the EGFR. The compounds are fused heterocyclic pyrimidine derivatives described at column 3, line 35 to column 5, line 6. The description of these compounds at column 3, line 35 to column 5, line 6 is incorporated herein by reference.
- [65] Barker, U.S. Patent 5,616,582 discloses quinazoline derivatives that have receptor tyrosine kinase inhibitory activity. The compounds disclosed in U.S. Patent 5,616,582 are incorporated herein by reference.
- [66] Fry et al., Science, 265: 1093-1095 (1994) discloses a compound having a structure that inhibits EGFR. The structure is shown in Figure 1. The compound shown in Figure 1 of the Fry et al. article is incorporated herein by reference.
- [67] Osherov et al., J. Biol. Chem., 268(15): 11,134-42 (1993) disclose tyrphostins that inhibit EGFR/HER1 and HER2. The compounds disclosed in the Osherov et al. article, and, in particular, those in Tables I, II, III, and IV are incorporated herein by reference.
- [68] Levitzki et al., U.S. Patent 5,196,446, discloses heteroarylethenediyl or heteroarylethenediylaryl compounds that inhibit EGFR. The compounds disclosed in U.S. Patent 5,196,446 from column 2, line 42 to column 3, line 40 are incorporated herein by reference.
- [69] Panek et al., J. Pharma. Exp. Thera., 283: 1433-1444 (1997) disclose a compound identified as PD166285 that inhibits the EGFR, PDGFR, and FGFR families of receptors. PD166285 is identified as 6-(2,6-dichlorophenyl)-2-(4-(2-diethylaminoethoxy)phenylamino)-8-methyl-8H-pyrido(2,3-d)pyrimidin-7-one having the structure shown in Figure 1 on page 1436. The compound described in Figure 1 on page 1436 of the Panek et al. article is incorporated herein by reference.
- [70] One example of a small molecule EGFR antagonist is IRESSA<sup>TM</sup> (ZD1939), which is a quinazoline derivative that functions as an ATP-mimetic to inhibit EGFR. See U.S. Patent No. 5,616,582 (Zeneca Limited); WO 96/33980 (Zeneca Limited) at p. 4; *see also*,

Rowinsky et al., Abstract 5 presented at the 37th Annual Meeting of ASCO, San Francisco, CA, 12-15 May 2001; Anido et al., Abstract 1712 presented at the 37th Annual Meeting of ASCO, San Francisco, CA, 12-15 May 2001.

[71] TARCEVA<sup>TM</sup> is another example of a small molecule EGFR antagonist. TARCEVA<sup>TM</sup> (OSI-774) is a 4-substituted phenylamino quinoxaline derivative [6,7-Bis(2-methoxy-ethoxy)-quinazolin-4-yl]- (3-ethynyl-phenyl)amine hydrochloride] EGFR inhibitor, which is described in WO 96/30347 (Pfizer Inc.) at, for example, page 2, line 12 through page 4, line 34 and page 19, lines 14-17. *See also* Moyer et al., *Cancer Res.*, 57: 4838-48 (1997); Pollack et al., *J. Pharmacol.*, 291: 739-48 (1999). TARCEVA<sup>TM</sup> may function by inhibiting phosphorylation of EGFR and its downstream PI3/Akt and MAP (mitogen activated protein) kinase signal transduction pathways resulting in p27-mediated cell-cycle arrest. *See* Hidalgo et al., Abstract 281 presented at the 37th Annual Meeting of ASCO, San Francisco, CA, 12-15 May 2001.

[72] Many other small molecules are known to inhibit EGFR. Some examples of small molecule EGFR antagonists are described in WO 91/116051, WO 96/30347, WO 96/33980, WO 97/27199 (Zeneca Limited), WO 97/30034 (Zeneca Limited), WO 97/42187 (Zeneca Limited), WO 97/49688 (Pfizer Inc.), WO 98/33798 (Warner Lambert Company), WO 00/18761 (American Cyanamid Company), and WO 00/31048 (Warner Lambert Company). Examples of specific small molecule EGFR antagonists include CI-1033, which is a quinoxaline (N-[4-(3-chloro-4-fluoro-phenylamino)-7-(3-morpholin-4-yl-propoxy)-quinazolin-6-yl]-acrylamide) inhibitor of tyrosine kinases, particularly EGFR and is described in WO 00/31048 (Warner-Lambert Company) at page 8, lines 22-6; PKI166, which is a pyrrolopyrimidine inhibitor of EGFR and is described in WO 97/27199 (Novartis AG) at pages 10-12; GW2016, which is an inhibitor of EGFR and HER2; and ExB569.

[73] Additional EGFR antagonists can easily be determined using well-known methods. The EGFR antagonists of the present invention inhibit the tyrosine kinase activity of EGFR, which generally involves phosphorylation events. Accordingly, phosphorylation assays are useful in determining antagonists useful in the context of the present invention. Some assays for tyrosine kinase activity are described in Panek et al. (1997) and Batley et al., *Life Sci.*, 62: 143-50 (1998). In addition, methods specific for detection of EGFR

expression can be utilized. These include immunohistochemistry (IHC) for detection of protein expression, fluorescence in situ hybridization (FISH) for detection of gene amplification, competitive radioligand binding assays, solid matrix blotting techniques, such as Northern and Southern blots, reverse transcriptase polymerase chain reaction (RT-PCR) and ELISA. *See, e.g.,* Grandis et al., *Cancer*, 78:1284-1292. (1996); Shimizu et al., *Japan J. Cancer Res.*, 85:567-571 (1994); Sauter et al., *Am. J. Path.*, 148:1047-1053 (1996); Collins, *Glia*, 15:289-296 (1995); Radinsky et al., *Clin. Cancer Res.*, 1:19-31 (1995); Petrides et al., *Cancer Res.*, 50:3934-3939 (1990); Hoffmann et al., *Anticancer Res.*, 17:4419-4426 (1997); Wikstrand et al., *Cancer Res.*, 55:3140-3148 (1995).

[74] *VEGF Receptor Antagonists*

[75] In one embodiment, the VEGF receptor antagonist binds specifically to an epitope on the extracellular domain of a VEGF receptor. The extracellular domain of a VEGF receptor is the ligand-binding domain. The ligand-binding domain may be found at either end of the receptor, but is normally found at the amino-terminal end.

[76] Some examples of VEGF receptors include the protein tyrosine kinase receptors referred to in the literature as flt-1 (VEGFR-1), KDR and flk-1 (VEGFR-2). Unless otherwise stated or clearly suggested otherwise by context, this specification will follow the customary literature nomenclature of VEGF receptors. KDR (VEGFR-2) will be referred to as the human form of a VEGF receptor having MW 180 kD (Terman et al., above). Flk-1 (VEGFR-2) will be referred to as the murine homolog of KDR (Matthews et al., above). Flt-1 (VEGFR-1) will be referred to as a form of VEGF receptor different from, but related to, the KDR/flk-1 (VEGFR-2) receptor. See Shibuya et al., above.

[77] Other VEGF receptors include those that can be cross-link labeled with VEGF, or that can be co-immunoprecipitated with KDR (VEGFR-2). Some known forms of these VEGF receptors have molecular weights of approximately 170 KD, 150 KD, 130-135 KD, 120-125 KD and 85 KD. *See, for example,* Quinn et al. *Proc. Nat'l Acad. Sci USA*, 90: 7533-7537 (1993); Scher et al., *J. Biol. Chem.*, 271: 5761-5767 (1996).

[78] The VEGF receptor is usually bound to a cell, such as an endothelial cell. The VEGF receptor may also be bound to a non-endothelial cell, such as a tumor cell. Alternatively, the VEGF receptor may be free from the cell, preferably in soluble form.

- [79] The antagonists of the present invention neutralize VEGF receptors. In this specification, neutralizing a receptor means inactivating the intrinsic kinase activity of the receptor to transduce a signal. A reliable assay for VEGF receptor neutralization is the inhibition of receptor phosphorylation.
- [80] The present invention is not limited by any particular mechanism of VEGF receptor neutralization. At the time of filing this application, the mechanism of VEGF receptor neutralization by antibodies was not well understood, and the mechanism followed by one antagonist is not necessarily the same as that followed by another antagonist. Some possible mechanisms include preventing binding of the VEGF ligand to the extracellular binding domain of the VEGF receptor, and preventing dimerization or oligomerization of receptors. Other mechanisms cannot, however, be ruled out.
- [81] A VEGF receptor (or VEGFR) antagonist, in the context of the present invention, is a biological molecule, small molecule, or any other substance that inhibits the VEGFR subfamily of receptors. By inhibition of activation of the VEGFR subfamily of receptors is meant any decrease in the activation of the VEGFR. That is, the prevention of activation need not completely stop activation of the VEGFR. Moreover, inhibition of VEGFR activation, as defined by the present invention, means inhibition of the VEGFR antagonist following interaction of the VEGFR antagonist and VEGFR. By association is meant sufficient physical or chemical interaction between the VEGFR antagonist and VEGFR that the receptor's tyrosine kinase activity is inhibited. One of skill in the art would appreciate that examples of such chemical interactions, which include association or bonding, are known in the art and include covalent bonding, ionic bonding, hydrogen bonding, etc. Accordingly, the VEGFR antagonists of the present invention inhibit the tyrosine kinase activity of the receptor, which prevents autophosphorylation of the receptor and phosphorylation of various other proteins involved in the VEGFR signaling pathways. Such pathways, which are involved in regulation of vasculogenesis and angiogenesis, include any of the following: the phospholipase C $\gamma$  (PLC $\gamma$ ) pathway or the phosphatidylinositol 3' kinase (PI3-K)/Akt and mitogen activated protein kinase (MAPK) pathway. *See, e.g., Larrivée et al., Int'l J. Mol. Med., 5: 447-56 (2000).*
- [82] The VEGFR subfamily of receptors is characterized by the presence of seven immunoglobulin-like loops in the extracellular domain, a single transmembrane region and

a split tyrosine kinase domain in the intracellular region (class III receptor tyrosine kinases). There are several known members of the VEGFR subfamily of receptors, examples of which include VEGFR-1, VEGFR-2, and VEGFR-3. It is generally believed that KDR (VEGFR-2) is the main VEGF signal transducer that results in endothelial cell proliferation, migration, differentiation, tube formation, increase of vascular permeability, and maintenance of vascular integrity. VEGFR-1 possesses a much weaker kinase activity, and is unable to generate a mitogenic response when stimulated by VEGF – although it binds to VEGF with an affinity that is approximately 10-fold higher than KDR (VEGFR-2). VEGFR-1 is also been implicated in VEGF and placenta growth factor (PlGF) induced migration of monocytes and macrophages and production of tissue factor.

[83] As is the case with EGFR described above, increased VEGFR activation can result from higher levels of ligand, VEGFR gene amplification, increased transcription of the receptor or mutations that cause unregulated receptor signaling.

[84] In one embodiment of the present invention, the VEGFR antagonist inhibits binding of VEGFR to its ligand. Binding of a ligand to an external, extracellular domain of VEGFR stimulates receptor dimerization, autophosphorylation of VEGFR, activation of the receptor's internal, cytoplasmic tyrosine kinase domain, and initiation of multiple signal transduction pathways involved in regulation of vasculogenesis and angiogenesis.

[85] Ligands for VEGFR include VEGF and its homologues PlGF, VEGF-B, VEGF-C, VEGF-D, and VEGF-E. For example, PlGF, which is a dimeric secreted factor that only binds VEGFR-1, is produced in large amounts by villous cytotrophoblast, syncytiotrophoblast and extravillous trophoblast and has close amino acid homology to VEGF. Three isoforms exist in humans, PlGF-1, PlGF-2, and PlGF-3. Studies with PlGF-deficient mice demonstrate that this growth factor is not involved in angiogenesis *per se*, but rather, specifically modulates the angiogenic and permeability effects of VEGF during pathological situations. Also, VEGF-D is closely related to VEGF-C by virtue of the presence of N- and C-terminal extensions that are not found in other VEGF family members. In adult human tissues, VEGF-D mRNA is most abundant in heart, lung, skeletal muscle, colon, and small intestine. Analyses of VEGF-D receptor specificity revealed that VEGF-D is a ligand for both VEGFR-2 (Flk1) and VEGFR-3 (Flt4) and can

activate these receptors; however, VEGF-D does not bind to VEGFR-1. In addition, VEGF-D is a mitogen for endothelial cells.

[86] In another embodiment of the present invention, the VEGFR antagonist binds VEGFR. It should be appreciated that the VEGFR antagonist can bind externally to the extracellular portion of VEGFR, which may or may not inhibit binding of the ligand, or internally to the tyrosine kinase domain. Examples of VEGFR antagonists that bind VEGFR include, without limitation, biological molecules, such as receptor ribozymes and antibodies (or functional equivalents thereof) specific for VEGFR, and synthetic kinase inhibitors that act directly on the cytoplasmic domain of VEGFR, such as small molecules. Preferably, the VEGFR antagonist of the present invention is a biological molecule and more preferably, an antibody, or functional equivalent thereof, specific for VEGFR, details of which are described in more detail below. Alternatively, the VEGFR antagonist of the present invention is a small molecule kinase inhibitor, details are described below.

[87] In one preferred embodiment, the VEGF receptor antagonist binds specifically to VEGFR-1. Particularly preferred are antigen-binding proteins that bind to the extracellular domain of VEGFR-1 and block binding by one or both of its ligands, VEGF and PlGF, and/or neutralize VEGF-induced or PlGF-induced activation of VEGFR-1. For example, mAb 6.12 is a scFv that binds to soluble and cell surface-expressed VEGFR-1. ScFv 6.12 comprises the V<sub>L</sub> and V<sub>H</sub> domains of mouse monoclonal antibody mAb 6.12. A hybridoma cell line producing mAb 6.12 has been deposited as ATCC number PTA-3344. The deposit was made under the provisions of the Budapest Treaty on the International Recognition of the Deposit of Microorganisms for the Purposes of Patent Procedure and the regulations thereunder (Budapest Treaty). This assures maintenance of a viable culture for 30 years from date of deposit. The organisms will be made available by ATCC under the terms of the Budapest Treaty, and subject to an agreement between Applicants and ATCC, which assures unrestricted availability upon issuance of the pertinent U.S. patent. Availability of the deposited strains is not to be construed as a license to practice the invention in contravention of the rights granted under the authority of any government in accordance with its patent laws.

- [88] Other preferred antibodies are described in the Examples, specifically Examples XII, XIII, and XIV, and in SEQ ID NO:1-83. Moreover, some of these preferred VEGFR antibody antagonists are also described in Lu et al., *Int'l J. Cancer*, 97: 393-399 (2002).
- [89] There also exist various hybridomas that produce VEGFR-2 antibodies. For example, a hybridoma cell line producing rat anti-mouse VEGFR-2 monoclonal antibody (DC101) was deposited as ATCC HB 11534; a hybridoma cell line (M25.18A1) producing mouse anti-mouse VEGFR-2 monoclonal antibody mAb 25 was deposited as ATCC HB 12152; a hybridoma cell line (M73.24) producing mouse anti-mouse VEGFR-2 monoclonal antibody mAb 73 was deposited as ATCC HB 12153.
- [90] In addition, there are various hybridomas that produce anti-VEGFR-1 antibodies include, but not limited to, hybridomas KM1730 (deposited as FERM BP-5697), KM1731 (deposited as FERM BP-5718), KM1732 (deposited as FERM BP-5698), KM1748 (deposited as FERM BP-5699), KM1750 (deposited as FERM BP-5700) disclosed in WO 98/22616, WO 99/59636, Australian accepted Application No. AU 1998 50666 B2, and Canadian Application No. CA 2328893.
- [91] Many other VEGFR antagonists are known in the art. Some examples of VEGFR antagonists are described in U.S. Application Nos. 07/813,593; 07/906,397; 07/946,507; 07/977,451; 08/055,269; 08/252,517; 08/601,891; 09/021,324; 09/208,786; and 09/919,408 (all to Lemischka et al.); U.S. Patent No. 5,840,301 (Rockwell et al.); U.S. Application Nos. 08/706,804; 08/866,969; 08/967,113; 09/047,807; 09/401,163; and 09/798,689 (all to Rockwell et al.); U.S. Application No. 09/540,770 (Witte et al.); and PCT/US01/06966 (Liao et al.). U.S. Patent No. 5,861,301 (Terman et al.), Terman et al. *Oncogene* 6: 1677-1683 (September 1991), WO 94/10202 (Ferrara et al.), and WO 95/21865 (Ludwig) disclose VEGFR antagonists and, specifically, anti-VEGFR-2 antibodies. In addition, PCT/US95/01678 (Kyowa Hakko), describes anti-VEGFR-2 antibodies. Anti-VEGFR antibodies are also described in U.S. Application No. 09/976,787 (Zhu et al.). U.S. Patent Nos. 6,177,401 (Ullrich et al.), 5,712,395 (App et al.), and 5,981,569 (App et al.) describe VEGFR antagonists that are organic molecules. In addition, bi-specific antibodies (BsAbs), which are antibodies that have two different antigen-binding specificities or sites, directed to KDR (VEGFR-2) and VEGFR-1 are known. *See, e.g.*, U.S. Application No. 09/865,198 (Zhu); 60/301,299 (Zhu).

- [92] Hennequin et al. in J. Med. Chem. 42, 5369-5389 (1999) disclose certain quinazolines, quinolines and cinnolines as being useful as VEGF receptor antagonists. See also Annie et al., Journal of Acquired Immune Deficiency Syndromes and Human Retrovirology 17, A41 (1998). The VEGF receptor antagonists disclosed in the Hennequin et al. article are incorporated herein by reference.
- [93] Additionally, App et al. (USPN: 5,849,742) discloses small molecule derivatives of quinazoline, quinoxiline, substituted aniline, isoxazoles, acrylonitrile and phenylacrylonitrile compounds which act as tyrosine kinase inhibitors. The small molecules described by Hennequin et al., Annie et al., and App et al. are included in the present invention as VEGF receptor antagonists.
- [94] Furthermore, assays for the determination of VEGFR antagonists are well known in the art and, therefore, alternate antagonists suitable for use in the present invention can be readily identified. The VEGFR antagonists of the present invention inhibit the tyrosine kinase activity of VEGFR, which generally involves phosphorylation events. Accordingly, phosphorylation assays are useful in determining VEGFR antagonists in the context of the present invention. Some assays for tyrosine kinase activity are described in Panek et al., J. Pharmacol. Exp. Thera., 283: 1433-44 (1997) and Batley et al., Life Sci., 62: 143-50 (1998). In addition, methods specific for detection of VEGFR expression can be utilized.
- [95] *Antibodies*
- [96] The antibodies of the present invention may be produced by methods known in the art. These methods include the immunological method described by Kohler and Milstein, Nature, 256: 495-497 (1975) and Campbell, Monoclonal Antibody Technology, The Production and Characterization of Rodent and Human Hybridomas, Burdon et al., Eds., Laboratory Techniques in Biochemistry and Molecular Biology, Volume 13, Elsevier Science Publishers, Amsterdam (1985); as well as by the recombinant DNA method described by Huse et al., Science, 246, 1275-1281 (1989).
- [97] The antibodies of the present invention can be monoclonal or polyclonal antibodies or any other suitable type of an antibody, such as a fragment or a derivative of an antibody, a single chain antibody (scFv) or a synthetic homolog of the antibody, provided

that the antibody has the same binding characteristics as, or that have binding characteristics comparable to, those of the whole antibody. As used herein, unless otherwise indicated or clear from the context, antibody domains, regions and fragments are accorded standard definitions as are well known in the art. *See, e.g.*, Abbas et al., Cellular and Molecular Immunology, W.B. Saunders Company, Philadelphia, PA (1991). Preferably, the antibodies of the subject invention are monoclonal antibodies.

- [98] Antibody fragments can be produced by cleaving a whole antibody, or by expressing DNA that encodes the fragment. Fragments of antibodies may be prepared by methods described by Lamoyi et al., J. Immunol. Methods, 56: 235-243 (1983) and by Parham, J. Immunol. 131: 2895-2902 (1983). Such fragments may contain one or both Fab fragments or the F(ab')<sub>2</sub> fragment. Such fragments may also contain single-chain fragment variable region antibodies, i.e. scFv, dibodies, or other antibody fragments. Methods of producing such functional equivalents are disclosed in PCT Application WO 93/21319, European Patent Application No. 239,400; PCT Application WO 89/09622; European Patent Application 338,745; and European Patent Application EP 332,424.
- [99] Single chain antibodies (scFv) are polypeptides that consist of the variable region of the heavy chain of the antibody linked to the variable region of the light chain with or without an interconnecting linker. Thus, the scFv comprises the entire antibody-combining site. These chains may be produced in bacteria, or in eukaryotic cells. An example of a single chain antibody is p1C11. (See Example IX below.) p1C11 was shown to block VEGF—KDR (VEGF—VEGFR-2) interaction and inhibit VEGF-stimulated receptor phosphorylation and mitogenesis of HUVEC. This scFv binds both soluble KDR (VEGFR-2) and cell surface-expressed KDR (VEGFR-2) on HUVEC. The sequence p1C11 of is shown as SEQ ID No: 21. The single chain antibodies described above can be built up into a chimerized or humanized antibody by methods known in the art; e.g., see example IX-3 below. One example of a chimerized scFv is chimerized p1C11, i.e., c-p1C11.
- [100] Preferably the antibody fragments contain all six complementarity-determining regions of the whole antibody, although fragments containing fewer than all of such regions, such as three, four or five CDRs, may also be functional. If the antibody fragment is too short to be immunogenic, it may be conjugated to a carrier molecule.

Some suitable carrier molecules include keyhole limpet hemocyanin and bovine serum albumen. Conjugation may be carried out by methods known in the art.

[101] Antibodies of the present invention also include those for which binding characteristics have been improved by direct mutation, methods of affinity maturation, phage display, or chain shuffling. Affinity and specificity may be modified or improved by mutating CDRs and screening for antigen binding sites having the desired characteristics (see, e.g., Yang et al., J. Mol. Bio., 254: 392-403 (1995)). CDRs are mutated in a variety of ways. One way is to randomize individual residues or combinations of residues so that in a population of otherwise identical antigen binding sites, all twenty amino acids are found at particular positions. Alternatively, mutations are induced over a range of CDR residues by error prone PCR methods (see, e.g., Hawkins et al., J. Mol. Bio., 226: 889-896 (1992)). Phage display vectors containing heavy and light chain variable region genes are propagated in mutator strains of *E. coli* (see, e.g., Low et al., J. Mol. Bio., 250: 359-368 (1996)). These methods of mutagenesis are illustrative of the many methods known to one of skill in the art.

[102] The antibodies of the present invention can also be chimeric antibodies having a variable region of an antibody of one species, for example, a mouse, and a constant region of an antibody of a different species, for example, a human. Alternatively, the antibodies of the present invention can be humanized antibodies having hypervariable or complementarity-determining regions (CDRs) of an antibody from one species, for example, a mouse, and framework variable regions and a constant region of a human antibody. Also alternatively, the antibodies of the present invention can be human antibodies having both a constant region and a variable region of a human antibody.

[103] Antibodies, and particularly monoclonal antibodies, can be produced by methods known in the art. Examples for production of antibodies include, but are not limited to, production in hybridoma cells and transformation of mammalian cells with DNA encoding the receptor antagonist. These methods are described in various publications, including the immunological method described by Kohler and Milstein, *Nature*, 256: 495-497 (1975) and Campbell in "Monoclonal Antibody Technology, The Production and Characterization of Rodent and Human Hybridomas" in Burdon et al., Eds., *Laboratory Techniques in Biochemistry and Molecular Biology*, Volume 13, Elsevier Science Publishers,

Amsterdam (1985); as well as by the recombinant DNA methods described by Huse et al. in *Science*, 246: 1275-1281 (1989).

- [104]       Equivalents of antibodies are also prepared by methods known in the art. For example, fragments of antibodies may be prepared enzymatically from whole antibodies. Preferably, equivalents of antibodies are prepared from DNA encoding such equivalents. DNA encoding fragments of antibodies may be prepared by deleting all but the desired portion of the DNA that encodes the full-length antibody. DNA encoding chimerized antibodies may be prepared by recombining DNA encoding human constant regions, derived substantially or exclusively from the corresponding human antibody regions, and DNA encoding variable regions, derived substantially or exclusively from the sequence of the variable region of a mammal other than a human. DNA encoding humanized antibodies may be prepared by recombining DNA encoding constant regions and variable regions other than the complementarity determining regions (CDRs), derived substantially or exclusively from the corresponding human antibody regions, and DNA encoding CDRs, derived substantially or exclusively from a mammal other than a human.
- [105]       Suitable sources of DNA molecules that encode fragments of antibodies include cells, such as hybridomas, that express the full-length antibody. The fragments may be used by themselves as antibody equivalents, or may be recombined into equivalents, as described above. The DNA deletions and recombinations described in this section may be carried out by known methods, such as those described in the published patent applications listed above in the section entitled "Functional Equivalents of Antibodies" and/or other standard recombinant DNA techniques, such as those described below.
- [106]       Preferred host cells for transformation of vectors and expression of the receptor antagonists of the present invention are mammalian cells, e.g., COS-7 cells, chinese hamster ovary (CHO) cells, and cell lines of lymphoid origin such as lymphoma, myeloma, or hybridoma cells. Other eukaryotic host, such as yeasts, can be alternatively used. For example, mouse fetal liver stromal cell line 2018 binds APTag-flk 1 and APTag-flk-2 fusion proteins, i.e., contains ligands of VEGFR-2 and flk-2 (ATCC, Manassas, VA, CRL 10907), human fetal spleen cell line Fsp 62891 contains flk-2 ligand (ATCC CRL 10935), and human stromal fetal thymus cell line, F.thy 62891, contains VEGFR-2 ligand (ATCC CRL 10936).

- [107] The transformed host cells are cultured by methods known in the art in a liquid medium containing assimilable sources of carbon (carbohydrates such as glucose or lactose), nitrogen (amino acids, peptides, proteins or their degradation products such as peptones, ammonium salts or the like), and inorganic salts (sulfates, phosphates and/or carbonates of sodium, potassium, magnesium and calcium). The medium furthermore contains, for example, growth-promoting substances, such as trace elements, for example iron, zinc, manganese and the like.
- [108] Where it is desired to express a gene construct in yeast, a suitable selection gene for use in yeast is the *trp1* gene present in the yeast plasmid YRp7. Stinchcomb et al. *Nature*, 282: 39 (1979); Kingsman et al., *Gene*, 7: 141 (1979). The *trp1* gene provides a selection marker for a mutant strain of yeast lacking the ability to grow in tryptophan, for example, ATCC No. 44076 or PEP4-1. Jones, *Genetics*, 85: 12 (1977). The presence of the *trp1* lesion in the yeast host cell genome then provides an effective environment for detecting transformation by growth in the absence of tryptophan. Similarly, *Leu2*-deficient yeast strains (ATCC 20,622 or 38,626) are complemented by known plasmids bearing the *Leu2* gene.
- [109] Also alternatively, the DNA encoding the receptor antagonist can be cloned into vectors derived from viruses such as adenovirus, adeno-associated virus, herpesvirus, retrovirus or lentivirus. Gene expression is controlled by inducible or uninducible regulatory sequences.
- [110] Briefly, a suitable source of cells containing nucleic acid molecules that express the desired DNA, such as an antibody, antibody equivalent or VEGF receptor, is selected. Total RNA is prepared by standard procedures from a suitable source. The total RNA is used to direct cDNA synthesis. Standard methods for isolating RNA and synthesizing cDNA are provided in standard manuals of molecular biology such as, for example, those described above.
- [111] The cDNA may be amplified by known methods. For example, the cDNA may be used as a template for amplification by polymerase chain reaction (PCR); see Saiki et al., *Science*, 239, 487 (1988) or Mullis et al., U.S. patent 4,683,195. The sequences of the oligonucleotide primers for the PCR amplification are derived from the known sequence

to be amplified. The oligonucleotides are synthesized by methods known in the art. Suitable methods include those described by Caruthers in Science 230, 281-285 (1985).

[112] A mixture of upstream and downstream oligonucleotides are used in the PCR amplification. The conditions are optimized for each particular primer pair according to standard procedures. The PCR product is analyzed, for example, by electrophoresis for cDNA having the correct size, corresponding to the sequence between the primers. Alternatively, the coding region may be amplified in two or more overlapping fragments. The overlapping fragments are designed to include a restriction site permitting the assembly of the intact cDNA from the fragments.

[113] In order to isolate the entire protein-coding regions for the VEGF receptors, for example, the upstream PCR oligonucleotide primer is complementary to the sequence at the 5' end, preferably encompassing the ATG start codon and at least 5-10 nucleotides upstream of the start codon. The downstream PCR oligonucleotide primer is complementary to the sequence at the 3' end of the desired DNA sequence. The desired DNA sequence preferably encodes the entire extracellular portion of the VEGF receptor, and optionally encodes all or part of the transmembrane region, and/or all or part of the intracellular region, including the stop codon.

[114] The DNA to be amplified, such as that encoding antibodies, antibody equivalents, or VEGF receptors, may also be replicated in a wide variety of cloning vectors in a wide variety of host cells. The host cell may be prokaryotic or eukaryotic.

[115] The vector into which the DNA is spliced may comprise segments of chromosomal, non-chromosomal and synthetic DNA sequences. Some suitable prokaryotic cloning vectors include plasmids from *E. coli*, such as *colE1*, *pCR1*, *pBR322*, *pMB9*, *pUC*, *pKSM*, and *RP4*. Prokaryotic vectors also include derivatives of phage DNA such as *M13* and other filamentous single-stranded DNA phages. A preferred vector for cloning nucleic acid encoding the VEGF receptor is the Baculovirus vector.

[116] The vector containing the DNA to be expressed is transfected into a suitable host cell. The host cell is maintained in an appropriate culture medium, and subjected to conditions under which the cells and the vector replicate. The vector may be recovered from the cell. The DNA to be expressed may be recovered from the vector.

- [117] The DNA to be expressed, such as that encoding antibodies, antibody equivalents, or receptors, may be inserted into a suitable expression vector and expressed in a suitable prokaryotic or eucaryotic host cell.
- [118] For example, the DNA inserted into a host cell may encode the entire extracellular portion of the VEGF receptor, or a soluble fragment of the extracellular portion of the VEGF receptor. The extracellular portion of the VEGF receptor encoded by the DNA is optionally attached at either, or both, the 5' end or the 3' end to additional amino acid sequences. The additional amino acid sequences may be attached to the VEGF receptor extracellular region in nature, such as the leader sequence, the transmembrane region and/or the intracellular region of the VEGF receptor. The additional amino acid sequences may also be sequences not attached to the VEGF receptor in nature. Preferably, such additional amino acid sequences serve a particular purpose, such as to improve expression levels, secretion, solubility, or immunogenicity.
- [119] Vectors for expressing proteins in bacteria, especially *E. coli*, are known. Such vectors include the PATH vectors described by Dieckmann and Tzagoloff in J. Biol. Chem. 260, 1513-1520 (1985). These vectors contain DNA sequences that encode anthranilate synthetase (TrpE) followed by a polylinker at the carboxy terminus. Other expression vector systems are based on beta-galactosidase (pEX); lambda P<sub>L</sub>; maltose binding protein (pMAL); and glutathione S-transferase (pGST) -see Gene 67, 31 (1988) and Peptide Research 3, 167 (1990).
- [120] Vectors useful in yeast are available. A suitable example is the 2 $\mu$  plasmid.
- [121] Suitable vectors for expression in mammalian cells are also known. Such vectors include well-known derivatives of SV-40, adenovirus, retrovirus-derived DNA sequences and shuttle vectors derived from combination of functional mammalian vectors, such as those described above, and functional plasmids and phage DNA.
- [122] Further eukaryotic expression vectors are known in the art (e.g., P.J. Southern and P. Berg, J. Mol. Appl. Genet., 1, 327-341 (1982); Subramani et al., Mol. Cell. Biol., 1: 854-864 (1981); Kaufmann and Sharp, "Amplification And Expression of Sequences Cotransfected with a Modular Dihydrofolate Reductase Complementary DNA Gene," J. Mol. Biol. 159, 601-621 (1982); Kaufmann and Sharp, Mol. Cell. Biol. 159, 601-664

(1982); Scahill et al., "Expression And Characterization Of The Product Of A Human Immune Interferon DNA Gene In Chinese Hamster Ovary Cells," Proc. Nat'l Acad. Sci. USA 80, 4654-4659 (1983); Urlaub and Chasin, Proc. Nat'l Acad. Sci. USA 77, 4216-4220, (1980).

[123] The expression vectors useful in the present invention contain at least one expression control sequence that is operatively linked to the DNA sequence or fragment to be expressed. The control sequence is inserted in the vector in order to control and to regulate the expression of the cloned DNA sequence. Examples of useful expression control sequences are the *lac* system, the *trp* system, the *tac* system, the *trc* system, major operator and promoter regions of phage lambda, the control region of fd coat protein, the glycolytic promoters of yeast, e.g., the promoter for 3-phosphoglycerate kinase, the promoters of yeast acid phosphatase, e.g., Pho5, the promoters of the yeast alpha-mating factors, and promoters derived from polyoma, adenovirus, retrovirus, and simian virus, e.g., the early and late promoters of SV40, and other sequences known to control the expression of genes of prokaryotic or eukaryotic cells and their viruses or combinations thereof.

[124] Vectors containing the control signals and DNA to be expressed, such as that encoding antibodies, antibody equivalents, or VEGF receptors, are inserted into a host cell for expression. Some useful expression host cells include well-known prokaryotic and eukaryotic cells. Some suitable prokaryotic hosts include, for example, *E. coli*, such as *E. coli* SG-936, *E. coli* HB 101, *E. coli* W3110, *E. coli* X1776, *E. coli* X2282, *E. coli* DHI, and *E. coli* MRC1, *Pseudomonas*, *Bacillus*, such as *Bacillus subtilis*, and *Streptomyces*. Suitable eukaryotic cells include yeast and other fungi, insect, animal cells, such as COS cells and CHO cells, human cells and plant cells in tissue culture.

[125] Following expression in a host cell maintained in a suitable medium, the polypeptide or peptide to be expressed, such as that encoding antibodies, antibody equivalents, or VEGF receptors, may be isolated from the medium, and purified by methods known in the art. If the polypeptide or peptide is not secreted into the culture medium, the host cells are lysed prior to isolation and purification.

[126] In addition, the antibodies of the invention may be prepared by immunizing a mammal with a soluble receptor. The soluble receptors may be used by themselves as

immunogens, or may be attached to a carrier protein or to other objects, such as beads, i.e. sepharose beads. After the mammal has produced antibodies, a mixture of antibody-producing cells, such as the splenocytes, is isolated. Monoclonal antibodies may be produced by isolating individual antibody-producing cells from the mixture and making the cells immortal by, for example, fusing them with tumor cells, such as myeloma cells. The resulting hybridomas are preserved in culture, and express monoclonal antibodies, which are harvested from the culture medium.

[127] The antibodies may also be prepared from receptors bound to the surface of cells that express the specific receptor of interest. The cell to which the receptors are bound may be a cell that naturally expresses the receptor, such as a vascular endothelial cell for VEGFR. Alternatively, the cell to which the receptor is bound may be a cell into which the DNA encoding the receptor has been transfected, such as 3T3 cells, which have been transfected with VEGFR.

[128] A receptor may be used as an immunogen to raise an antibody of the invention. The receptor peptide may be obtained from natural sources, such as from cells that express the receptors. For example, the VEGF receptor peptide may be obtained from vascular endothelial cells. Alternatively, synthetic receptor peptides may be prepared using commercially available machines. In such an embodiment, the VEGF receptor amino acid sequence can be provided by, for example, Shibuya et al., *Oncogene* 5, 519-524 (1990) for flt-1 (VEGFR-1); PCT/US92/01300 and Terman et al., *Oncogene* 6:1677-1683 (1991) for KDR (VEGFR-2); and Matthews et al. *Proc. Nat'l Acad. Sci. USA*, 88:9026-9030 (1991) for flk-1.

[129] As a further alternative, DNA encoding a receptor, such as a cDNA or a fragment thereof, may be cloned and expressed and the resulting polypeptide recovered and used as an immunogen to raise an antibody of the invention. For example, in order to prepare the VEGF receptors against which the antibodies are made, nucleic acid molecules that encode the VEGF receptors of the invention, or portions thereof, especially the extracellular portions thereof, may be inserted into known vectors for expression in host cells using standard recombinant DNA techniques, such as those described below. Suitable sources of such nucleic acid molecules include cells that express VEGF receptors, i.e. vascular endothelial cells.

- [130] The antibody may be prepared in any mammal; suitable mammals other than a human include, for example, a rabbit, rat, mouse, horse, goat, or primate. Mice are preferred. The antibody may be a member of one of the following immunoglobulin classes: IgG, IgM, IgA, IgD, or IgE, and the subclasses thereof, and preferably is an IgG1 antibody. The antibodies of the invention and their functional equivalents may be or may combine members of any of the immunoglobulin classes.
- [131] *Non-Antibody VEGFR Antagonists*
- [132] In addition to the antibodies, or functional equivalents of antibodies, discussed above, the receptor antagonists useful in the present invention may also be other biological and small molecules, especially in connection with the treatments described above.
- [133] Biological molecules include all lipids and polymers of monosaccharides, amino acids and nucleotides having a molecular weight greater than 450. Thus, biological molecules include, for example, oligosaccharides and polysaccharides; oligopeptides, polypeptides, peptides, and proteins; and oligonucleotides and polynucleotides. Oligonucleotides and polynucleotides include, for example, DNA and RNA.
- [134] Biological molecules further include derivatives of any of the molecules described above. For example, derivatives of biological molecules include lipid and glycosylation derivatives of oligopeptides, polypeptides, peptides and proteins. Derivatives of biological molecules further include lipid derivatives of oligosaccharides and polysaccharides, for example, lipopolysaccharides.
- [135] Any molecule that is not a biological molecule is considered in this specification to be a small molecule. Small molecules of the present invention are entities having carbon and hydrogen atoms, as well as heteroatoms, which include, but are not limited to, nitrogen, sulfur, oxygen, and phosphorus. Atoms in a small molecule are linked together via covalent and ionic bonds; the former is typical for small organic compounds, e.g., small molecule tyrosine kinase inhibitors such as Iressa<sup>TM</sup> and Tarceva<sup>TM</sup> and the latter is typical of small inorganic compounds. The arrangement of atoms in a small organic molecule may represent a chain, e.g., a carbon-carbon chain or carbon-heteroatom chain, or ring containing carbon atoms, e.g., benzene, or a combination of carbon and heteroatoms, i.e., heterocycles, for example, a pyrimidine or quinazoline. A combination

of one or more chains in a small organic molecule attached to a ring system constitutes a substituted ring system and fusion of two rings constitutes a fused polycyclic system, which can be referred to as simply a polycyclic system, an example of which is the parent scaffold of Iressa.<sup>TM</sup> Small molecules include both compounds found in nature, such as hormones, neurotransmitters, nucleotides, amino acids, sugars, lipids and their derivatives, and those compounds made synthetically, either by traditional organic synthesis, bio-mediated synthesis, or a combination thereof. *See, e.g.,* Ganesan, Drug Discov. Today, 7(1): 47-55 (Jan. 2002); Lou, Drug Discov. Today, 6(24): 1288-1294 (Dec. 2001).

- [136] Some examples of small molecules include organic compounds, organometallic compounds, salts of organic and organometallic compounds, saccharides, amino acids, nucleosides and nucleotides. It is emphasized that small molecules can have any molecular weight. They are merely called small molecules because they typically have molecular weights less than 450. Small molecules include compounds that are found in nature as well as synthetic compounds.
- [137] The administration of small molecule and biological drugs to human patients is accomplished by methods known in the art. Examples of such methods for small molecules are described in Spada, U.S. Patent 5,646,153 at column 57, line 47 to column 59, line 67. This description of administering small molecules is incorporated herein by reference.
- [138] *Neutralizing VEGF Activation of VEGF Receptors*
- [139] Neutralization of activation of a VEGF receptor in a sample of endothelial or non-endothelial cells, such as tumor cells, may be performed *in vitro* or *in vivo*. Neutralizing VEGF activation of a VEGF receptor in a sample of VEGF-receptor expressing cells comprises contacting the cells with an antagonist, e.g., an antibody, of the invention. The cells are contacted *in vitro* with the antagonist, e.g., the antibody, before, simultaneously with, or after, adding VEGF to the cell sample.
- [140] *In vivo*, an antagonist, e.g., an antibody, of the invention is contacted with a VEGF receptor by administration to a mammal. Methods of administration to a mammal include, for example, oral, intravenous, intraperitoneal, subcutaneous, or intramuscular administration.

- [141] This *in vivo* neutralization method is useful for inhibiting angiogenesis in a mammal. Angiogenesis inhibition is a useful therapeutic method, such as for preventing or inhibiting angiogenesis associated with pathological conditions such as tumor growth. Accordingly, the antagonists, e.g., the antibodies, of the invention are anti-angiogenic and anti-tumor immunotherapeutic agents.
- [142] The word mammal means any mammal. Some examples of mammals include pet animals, such as dogs and cats; farm animals, such as pigs, cattle, sheep, and goats; laboratory animals, such as mice and rats; primates, such as monkeys, apes, and chimpanzees; and humans.
- [143] VEGF receptors are found on some non-endothelial cells, such as tumor cells, indicating the unexpected presence of an autocrine and/or paracrine loop in these cells. The antagonists, e.g., the antibodies, of this invention are useful in neutralizing activity of VEGF receptors on such cells, thereby blocking the autocrine and/or paracrine loop, and inhibiting tumor growth.
- [144] The methods of inhibiting angiogenesis and of inhibiting pathological conditions such as tumor growth in a mammal comprise administering an effective amount of any one of the invention's antagonists, e.g., antibodies, including any of the functional equivalents thereof, systemically to a mammal, or directly to a tumor within the mammal. The mammal is preferably human. This method is effective for treating subjects with both solid tumors, preferably highly vascular tumors, and non-solid tumors.
- [145] The inhibition or reduction of tumor growth includes the prevention or inhibition of the progression of a tumor, including cancerous and noncancerous tumors. The progression of a tumor includes the invasiveness, metastasis, recurrence and increase in size of the tumor. The inhibition or reduction of tumor growth also includes the destruction of a tumor.
- [146] All types of tumors may be treated by the methods of the present invention. The tumors may be solid or non-solid.
- [147] Some examples of solid tumors that can be treated with the antagonists of the present invention include carcinomas, sarcomas, blastomas or gliomas. Some examples of

such tumors include epidermoid tumors, squamous tumors, such as head and neck tumors, colorectal tumors, prostate tumors, breast tumors, lung tumors, including small cell and non-small cell lung tumors, pancreatic tumors, thyroid tumors, ovarian tumors, and liver tumors. Other examples include Kaposi's sarcoma, CNS neoplasms, neuroblastomas, capillary hemangioblastomas, meningiomas and cerebral metastases, melanoma, gastrointestinal and renal carcinomas and sarcomas, rhabdomyosarcoma, glioblastoma, preferably glioblastoma multiforme, and leiomyosarcoma. Examples of vascularized skin cancers for which the antagonists of this invention are effective include squamous cell carcinoma, basal cell carcinoma and skin cancers that can be treated by suppressing the growth of malignant keratinocytes, such as human malignant keratinocytes.

- [148] Some examples of non-solid tumors include leukemias, multiple myelomas and lymphomas. Some examples of leukemias include acute myelocytic leukemia (AML), chronic myelocytic leukemia (CML), acute lymphocytic leukemia (ALL), chronic lymphocytic leukemia (CLL), erythrocytic leukemia or monocytic leukemia. Some examples of lymphomas include lymphomas associated with Hodgkin's disease and Non-Hodgkin's disease.
- [149] Experimental results described later demonstrate that antibodies of the invention specifically block VEGF induced phosphorylation of a mouse extracellular flk-1 (VEGFR-2) (VEGFR-2)/intracellular fms chimeric receptor expressed in transfected 3T3 cells. The antibodies had no effect on a fully stimulated chimeric extracellular fms/intracellular FLK-2 receptor by CSF-1. *In vivo* studies also described below show that the antibodies were able to significantly inhibit tumor growth in nude mice.
- [150] A cocktail of VEGF receptor antagonists, e.g., monoclonal antibodies, provides an especially efficient treatment for inhibiting the growth of tumor cells. The cocktail may include as few as 2, 3 or 4 antibodies, and as many as 6, 8 or 10 antibodies.
- [151] Preventing or inhibiting angiogenesis is also useful to treat non-neoplastic pathologic conditions characterized by excessive angiogenesis, such as neovascular glaucoma, proliferative retinopathy including proliferative diabetic retinopathy, arthritis, macular degeneration, hemangiomas, angiofibromas, and psoriasis.

**[152]** *Using The Antagonists of The Invention To Isolate and Purify the VEGF Receptor*

**[153]** The antagonists of the present invention may be used to isolate and purify the VEGF receptor using conventional methods such as affinity chromatography (Dean et al., *Affinity Chromatography: A Practical Approach*, IRL Press, Arlington, VA (1985)). Other methods well known in the art include magnetic separation with antibody-coated magnetic beads, "panning" with an antibody attached to a solid matrix, and flow cytometry.

**[154]** The source of the VEGF receptor is typically vascular cells, and especially vascular endothelial cells, that express the VEGF receptor. Suitable sources of vascular endothelial cells are blood vessels, such as umbilical cord blood cells, especially, human umbilical cord vascular endothelial cells (HUVEC).

**[155]** The VEGF receptors may be used as starting material to produce other materials, such as antigens for making additional monoclonal and polyclonal antibodies that recognize and bind to the VEGF receptor or other antigens on the surface of VEGF-expressing cells.

**[156]** *Using the Antagonists of the Invention to Isolate and Purify flk-1 (VEGFR-2) Positive Tumor Cells*

**[157]** The antagonists of the present invention may be used to isolate and purify flk-1 (VEGFR-2) (*VEGFR-2*) positive tumor cells, i.e., tumor cells expressing the flk-1 (VEGFR-2) receptor, using conventional methods such as affinity chromatography (Dean, P.D.G. et al., *Affinity Chromatography: A Practical Approach*, IRL Press, Arlington, VA (1985)). Other methods well known in the art include magnetic separation with antibody-coated magnetic beads, cytotoxic agents, such as complement, conjugated to the antibody, "panning" with an antibody attached to a solid matrix, and flow cytometry.

**[158]** *Monitoring Levels of VEGF and VEGF Receptors In Vitro or In Vivo*

**[159]** The antagonists, e.g., antibodies, of the invention may be used to monitor levels of VEGF or VEGF receptors *in vitro* or *in vivo* in biological samples using standard assays

and methods known in the art. Some examples of biological samples include bodily fluids, such as blood. Standard assays involve, for example, labeling the antibodies and conducting standard immunoassays, such as radioimmunoassays, as is well known in the art.

[160] Standard recombinant DNA techniques useful in carrying out the present invention are described in Sambrook et al., "Molecular Cloning," Second Edition, Cold Spring Harbor Laboratory Press (1987) and by Ausubel et al. (Eds) "Current Protocols in Molecular Biology," Green Publishing Associates/ Wiley-Interscience, New York (1990).

[161] *Administration*

[162] The present receptor antagonists can be administered for therapeutic treatments to a patient suffering from a tumor in an amount sufficient to prevent, inhibit, or reduce the progression of the tumor, e.g., the growth, invasiveness, metastases and/or recurrence of the tumor. An amount adequate to accomplish this is defined as a therapeutically effective dose. Amounts effective for this use will depend upon the severity of the disease and the general state of the patient's own immune system. Dosing schedules will also vary with the disease state and status of the patient, and will typically range from a single bolus dosage or continuous infusion to multiple administrations per day (e.g., every 4-6 hours), or as indicated by the treating physician and the patient's condition. It should be noted, however, that the present invention is not limited to any particular dose.

[163] The present invention can be used to treat any suitable tumor, including, for example, tumors of the breast, heart, lung, small intestine, colon, spleen, kidney, bladder, head and neck, ovary, prostate, brain, pancreas, skin, bone, bone marrow, blood, thymus, uterus, testicles, cervix or liver. Preferably, the present methods are used when the tumor is a tumor of the colon or when the tumor is a non-small cell lung carcinoma (NSCLC).

[164] Moreover, the tumors of the present invention preferably overexpress EGFR. Enhanced expression of EGFR has been detected in a significant percentage of many human tumors; for example, head and neck (80-100 %), colorectal (25-77 %), pancreatic (30-50 %), lung (40-80 %), esophageal (43-89 %), renal cell (50-90 %), prostate (65 %), bladder (31-48 %), cervical/uterus (90 %), ovarian (35-70 %) and breast (14-91 %).

Expression of EGFR has also been demonstrated to be an indicator of poor prognosis, decreased survival, and increased metastases in certain tumor types.

[165] Furthermore, the tumors of the present invention preferably have aberrant expression or signaling of VEGFR. Enhanced signaling by VEGFR has been observed in many different human cancers. High levels of VEGFR-2 are expressed by endothelial cells that infiltrate gliomas (Plate et al., (1992) *Nature* 359:845-848). VEGFR-2 levels are specifically upregulated by VEGF produced by human glioblastomas (Plate et al. (1993) *Cancer Res.* 53:5822-5827). The finding of high levels of VEGFR-2 expression in glioblastoma associated endothelial cells (GAEC) indicates that receptor activity is probably induced during tumor formation since VEGFR-2 transcripts are barely detectable in normal brain endothelial cells. This upregulation is confined to the vascular endothelial cells in close proximity to the tumor.

[166] The present invention is useful for inhibition or reduction of tumor growth. By inhibition or reduction of tumor growth is meant prevention, inhibition, or reduction of the progression of the tumor, e.g, the growth, invasiveness, metastases and/or recurrence of the tumor. In addition, the present invention also can be useful in treating an angiogenic condition, such as atherosclerosis, arthritis, macular degeneration and psoriasis. The identification of those patients that have conditions for which the present invention is useful is well within the ability and knowledge of one skilled in the art.

[167] In the present invention, any suitable method or route can be used to administer the EGFR antagonist and/or the VEGFR antagonist, for example, oral, intravenous, intraperitoneal, subcutaneous, or intramuscular administration. The dose of antagonist administered depends on numerous factors, including, for example, the type of antagonists, the type and severity tumor being treated and the route of administration of the antagonists. It should be emphasized, however, that the present invention is not limited to any particular method or route of administration.

[168] In one alternative embodiment, the EGFR antagonist and the VEGFR antagonist can be administered in combination with one or more antineoplastic agents. *See, e.g.,* U.S. Patent No. 6,217,866 (Schlessinger et al.) (Anti-EGFR antibodies in combination with antineoplastic agents); U.S. Application No. 09/312,286 (Waksal et al.) (Anti-EGFR antibodies in combination with radiation). Any suitable antineoplastic agent can be used,

such as a chemotherapeutic agent or radiation. Examples of chemotherapeutic agents include, but are not limited to, cisplatin, doxorubicin, paclitaxel, irinotecan (CPT-11), topotecan or a combination thereof. When the antineoplastic agent is radiation, the source of the radiation can be either external (external beam radiation therapy – EBRT) or internal (brachytherapy – BT) to the patient being treated. The dose of antineoplastic agent administered depends on numerous factors, including, for example, the type of agent, the type and severity tumor being treated and the route of administration of the agent. It should be emphasized, however, that the present invention is not limited to any particular dose.

[169] In an additional alternative embodiment, the EGFR antagonist and the VEGFR antagonist can be administered in combination with one or more suitable adjuvants, such as, for example, cytokines (IL-10 and IL-13, for example) or other immune stimulators. *See, e.g., Larrivée et al., supra.* It should be appreciated, however, that administration of only an EGFR antagonist and a VEGFR antagonist is sufficient to prevent, inhibit, or reduce the progression of the tumor in a therapeutically effective manner.

[170] In addition, the EGFR antagonist and/or the VEGFR antagonist can be administered as a ligand conjugate, which binds specifically to the receptor and deliver a toxic, lethal payload following ligand-toxin internalization. Conjugates between toxins and the receptors were designed with the aim of developing toxic agents specific for EGFR- or VEGFR-overexpressing tumor cells while minimizing nonspecific toxicity. For example, a conjugate composed of EGF and *Pseudomonas* endotoxin (PE) was shown to be toxic toward EGFR-expressing HeLa cells in vitro. Various agents, including thioridazine and adenovirus, can enhance cellular uptake of the conjugate, as well as increase the cytotoxicity of the conjugate.

[171] It is understood that the EGFR and/or VEGFR antagonists of the invention, where used in a mammal for the purpose of prophylaxis or treatment, will be administered in the form of a composition additionally comprising a pharmaceutically acceptable carrier. Suitable pharmaceutically acceptable carriers include, for example, one or more of water, saline, phosphate buffered saline, dextrose, glycerol, ethanol and the like, as well as combinations thereof. Pharmaceutically acceptable carriers may further comprise minor amounts of auxiliary substances such as wetting or emulsifying agents, preservatives or

buffers, which enhance the shelf life or effectiveness of the binding proteins. The compositions of the injection may, as is well known in the art, be formulated so as to provide quick, sustained or delayed release of the active ingredient after administration to the mammal.

[172] The EGFR antagonists and/or VEGFR antagonists of this invention may be in a variety of forms. These include, for example, solid, semi-solid and liquid dosage forms, such as tablets, pills, powders, liquid solutions, dispersions or suspensions, liposomes, suppositories, injectable and infusible solutions. The preferred form depends on the intended mode of administration and therapeutic application.

[173] Such antagonists can be prepared in a manner well known in the pharmaceutical art. In making the composition the active ingredient will usually be mixed with a carrier, or diluted by a carrier, and/or enclosed within a carrier which may, for example, be in the form of a capsule, sachet, paper or other container. When the carrier serves as a diluent, it may be a solid, semi-solid, or liquid material, which acts as a vehicle, excipient or medium for the active ingredient. Thus, the composition may be in the form of tablets, lozenges, sachets, cachets, elixirs, suspensions, aerosols (as a solid or in a liquid medium), ointments containing for example up to 10% by weight of the active compound, soft and hard gelatin capsules, suppositories, injection solutions, suspensions, sterile packaged powders and as a topical patch.

[174] *Kits for Inhibition of Tumor Growth*

[175] The present invention also includes kits for inhibiting tumor growth comprising a therapeutically effective amount of an EGFR antagonist and a therapeutically effective amount of a VEGFR antagonist. The EGFR or VEGFR antagonist of the present kits can be any suitable antagonist, examples have been described above. Preferably, the EGFR antagonist of the kit comprises an antibody, or functional equivalent thereof, specific for EGFR. Alternatively, and also preferably, the EGFR antagonist of the kit comprises a small molecule specific for EGFR. The VEGFR antagonist of the kit preferably comprises an antibody, or functional equivalent thereof, specific for VEGFR. Alternatively, the VEGFR antagonist of the kit preferably comprises a small molecule specific for VEGFR. In addition, the kits of the present invention can further comprise an antineoplastic agent. Examples of suitable antineoplastic agents in the context of the present invention have

been described herein. The kits of the present invention can further comprise an adjuvant, examples have also been described above.

[176] Accordingly, the present receptor antagonists thus can be used *in vivo* and *in vitro* for investigative, diagnostic, prophylactic, or treatment methods, which are well known in the art. Of course, it is to be understood and expected that variations in the principles of invention herein disclosed can be made by one skilled in the art and it is intended that such modifications are to be included within the scope of the present invention.

[177] All references mentioned herein are incorporated in their entirety.

### EXAMPLES

[178] The Examples that follow are set forth to aid in understanding the invention but are not intended to, and should not be construed to, limit its scope in any way. The Examples do not include detailed descriptions of conventional methods, such as those employed in the construction of vectors and plasmids, the insertion of genes encoding polypeptides into such vectors and plasmids, or the introduction of plasmids into host cells. Such methods are well known to those of ordinary skill in the art and are described in numerous publications including Sambrook, J., Fritsch, E.F. and Maniatis, T. (1989) *Molecular Cloning: A Laboratory Manual*, 2nd edition, Cold Spring Harbor Laboratory Press.

#### [179] *EXAMPLE I. CELL LINES AND MEDIA*

[180] NIH 3T3 cells were obtained from the American Type Culture Collection (Rockville MD). The C441 cell line was constructed by transfecting 3T3 cells with the chimeric receptor mouse flk-1 (VEGFR-2)/human fms. 10A2 is a 3T3 transfectant containing the chimeric receptor human fms/mouse *FLK-2*, the isolation and characterization of which has been described (Dosil et al., *Mol. Cell. Biol.* 13:6572-6585 (1993)). Cells were routinely maintained in Dulbecco's modified Eagle's medium (DME) supplemented with 10% calf serum (CS), 1 mM L-glutamine, antibiotics, and 600 ug/ml G418 (Geneticin; Sigma, St Louis MO).

[181] A glioblastoma cell line, GBM-18, was maintained in DME supplemented with 5% calf serum, 1mM L-glutamine, and antibiotics.

[182] A stable 3T3 line secreting the soluble chimeric protein, mouse flk-1 (VEGFR-2):SEAPs (secretory alkaline phosphatase), was generated and maintained in DMEM and 10% calf serum. Conditioned media was collected. Soluble flk-1 (VEGFR-2):SEAP is isolated from the conditioned media.

[183] *EXAMPLE II. ISOLATION OF MONOCLONAL ANTIBODIES*

[184] *Example II-1. Rat anti mouse flk-1 (VEGFR-2) monoclonal antibody DC-101 (IgG1)*

[185] Lewis rats (Charles River Labs) were hyperimmunized with an immune complex consisting of the mouse flk-1:SEAPs soluble receptor, a rabbit anti-alkaline phosphatase polyclonal antibody and Protein-G sepharose beads. The animals received 7 intraperitoneal injections of this complex spread over three months (at days 0, 14, 21, 28, 49, 63, 77). At various times, the animals were bled from the tail vein and immune sera screened by ELISA for high titer binding to mflk-1 (VEGFR-2): SEAPs. Five days after the final injection, rats were sacrificed and the spleens aseptically removed. Splenocytes were washed, counted, and fused at a 2:1 ratio with the murine myeloma cell line NS1. Hybridomas were selected in HAT medium and colonies screened by ELISA for specific binding to mflk-1 (VEGFR-2):SEAPs but not the SEAPs protein. A number of positive hybridomas were expanded and cloned three times by limiting dilution. One subclone, designated DC-101, was further characterized.

[186] *Example II-2. Mouse anti mouse flk-1 (VEGFR-2) monoclonal antibodies Mab 25 and Mab 73*

[187] Murine anti-flk-1 (VEGFR-2) monoclonal antibodies (Mabs) were produced using a similar protocol as that employed for DC-101. Briefly, mice were injected with a complex of flk-1 (VEGFR-2)/SEAP soluble receptor bound to either an anti-SEAP-Protein/A Sepharose complex or wheat germ agglutinin Sepharose from conditioned medium of transfected NIH 3T3 cell. Mice were hyperimmunized at periodic intervals over a 6 month period. Immune splenocytes were pooled and fused with the murine myeloma cell line, NSI. Hybridomas were selected in HAT medium and following incubation, colonies were screened for mouse Mab production. Unlike the protocol employed for DC-101, positive supernatants were initially screened for binding to the flk-1 (VEGFR-2)/fms receptor captured from C441 cell lysates on ELISA plates coated with a

peptide generated polyclonal antibody against the C-terminal region of *fms*. Reactive Mabs were then assayed by ELISA for binding to intact C441 cells and to purified *flk-1* (VEGFR-2)/SEAP versus SEAP alone. The supernatants from hybridomas showing binding to C441 and reactivity with *flk-1* (VEGFR-2)/SEAP but not SEAP were expanded, grown in ascites, and purified (EZ-PREP, Pharmacia). Purified Mabs were subjected to assays on C441 cells to determine their cell surface binding by FACS and their ability to inhibit VEGF induced activation of *flk-1* (VEGFR-2)/*fms* in phosphorylation assays. The results of these studies led to the cloning of Mabs 25 and 73 (isotype IgG1) for further characterization based on their capabilities to bind specifically to *flk-1* (VEGFR-2) and block receptor activation at levels comparable to that observed for DC-101.

[188] *EXAMPLE III. ASSAYS*

[189] *Example III-1. ELISA Methods*

[190] Antibodies were screened by a solid state ELISA in which the binding characteristics of the various mAbs to *flk-1* (VEGFR-2):SEAP and SEAP protein were compared. Microtiter plates were coated with 50-100 ng/well of either *flk-1*:SEAP or AP in pH9.6 carbonate buffer overnight at 4°C. Plates were blocked with phosphate buffered saline supplemented with 10% new born calf serum (NB10) for one hour at 37°C. Hybridoma supernatants or purified antibodies were added to the plates for two hours at 37°C followed by goat anti-rat IgG conjugated to horseradish peroxidase (Tago) added for an additional hour at 37°C. After extensive washing, TMB (Kirkegaard and Perry, Gaithersburg MD) plus hydrogen peroxide was added as the chromogen and the plates read at 450 nm in an ELISA reader.

[191]        *Example III-2. Isotyping*

[192]        Isotyping of the various monoclonal antibodies was done as previously described (Songsakphisarn, R. and Goldstein, N.I., Hybridoma 12: 343-348, 1993) using rat isotype specific reagents (Zymed Labs, South San Francisco CA).

[193]        *Example III-3. Phosphorylation, Immunoprecipitation and Immunoblot Assays*

[194]        The phosphorylation assays and Western blot analysis with C441 and 10A2 cells were performed as previously described (Tessler et al., 1994) with some modifications. Briefly, cells were grown to 90% confluency in DME-10% CS and then serum starved in DME-0.5% CS for 24 hours prior to experimentation. HUVEC cells were grown to subconfluence in EGM basal media. For neutralization assays, cells were stimulated with various concentrations of the appropriate ligand under serum free conditions (DME -0.1% BSA) in the presence and absence of mAb DC-101 for 15 minutes at room temperature. The ligands, VEGF and CSF-1, were assayed at concentrations of 10-80 ng/ml and 20-40 ng/ml, respectively. Monoclonal antibodies were assayed at concentrations ranging from 0.5 µg/ml to 10 µg/ml. To evaluate the effects of mAb DC-101 on the VEGF induced activation of the flk-1 (VEGFR-2)-fms receptor, antibody was either added simultaneously (competitive inhibition) or prebound to cells for 15 minutes at room temperature prior to the addition of ligand. Cells incubated in serum free medium in the absence and presence of DC-101 served as controls for receptor autophosphorylation in the absence of ligand and the presence of antibody, respectively. A control cell line expressing the fms/FLK-2 chimeric receptor (10A2) was starved and stimulated with 20 and 40 ng/ml CSF-1 and assayed in the presence and absence of 5 µg/ml DC-101.

[195]        Following stimulation, monolayers were washed with ice cold PBS containing 1mM sodium orthovanadate. Cells were then lysed in lysis buffer (20 mM Tris-HCl, pH 7.4, 1% Triton X-100, 137mM NaCl, 10% glycerol, 10 mM EDTA, 2mM sodium orthovanadate, 100 mM NaF, 100mM sodium pyrophosphate, 5mM Pefabloc (Boehringer Mannheim Biochemicals, Indianapolis IN), 100 µg aprotinin and 100 µg/ml leupeptin) and centrifuged at 14000 x g for 10 minutes. Protein was immunoprecipitated from cleared lysates of transfected cells using polyclonal antibodies generated to peptides corresponding to the C-terminal region of the human fms receptor (Tessler et al., J. Biol. Chem. 269, 12456-12461, 1994) or the murine FLK-2 interkinase domain (Small et al.,

Proc. Nat'l Acad. Sci. USA, 91, 459-463, 1994) coupled to Protein A Sepharose beads. Where indicated, immunoprecipitations with DC-101 or irrelevant rat IgG were performed with 10 µg of antibody coupled to Protein G beads. The beads were then washed once with 0.2% Triton X-100, 10 mM Tris-HCl pH8.0, 150 mM NaCl, 2mM EDTA (Buffer A), twice with Buffer A containing 500 mM NaCl and twice with Tris-HCl, pH 8.0. Drained beads were mixed with 30 µl in 2x SDS loading buffer and subjected to SDS PAGE in 4-12% gradient gels (Novex, San Diego CA). After electrophoresis, proteins were blotted to nitrocellulose filters for analysis. Filters were blocked overnight in blocking buffer (50 mM Tris-HCl, pH7.4, 150 mM NaCl (TBS) containing 5% bovine serum albumin and 10% nonfat dried milk (Biorad, CA). To detect phosphorylated receptor, blots were probed with a monoclonal antibody directed to phosphotyrosine (UBI, Lake Placid, NY) in blocking buffer for 1 hour at room temperature. Blots were then washed extensively with 0.5 x TBS containing 0.1% Tween-20 (TBS-T) and incubated with goat anti-mouse Ig conjugated to horseradish peroxidase (Amersham). Blots were washed with TBS and incubated for 1 minute with a chemiluminescence reagent (ECL, Amersham). Anti-phosphotyrosine reacting with phosphorylated proteins was detected by exposure to a high performance luminescence detection film (Hyperfilm-ECL, Amersham) for 0.5 to 10 minutes.

[196] To detect flk-1 (VEGFR-2)/fms in C441 cells receptor levels, blots were stripped according to manufacturer's protocols (Amersham) and reprobed with the anti-fms rabbit polyclonal antibody.

[197] *Example III-4. Flow Cytometer Binding Assays*

[198] C441 cells were grown to near confluency in 10 cm plates. Cells were removed with a non-enzymatic dissociation buffer (Sigma), washed in cold serum free medium and resuspended in Hanks balanced salt solution supplemented with 1% BSA (HBSS-BSA) at a concentration of 1 million cells per tube. Monoclonal Ab DC-101 or an isotype matched irrelevant antibody anti *FLK-2* 23H7 was added at 10 µg per tube for 60 minutes on ice. After washing, 5µl of goat anti-mouse IgG conjugated to FITC (TAGO) was added for an additional 30 minutes on ice. Cells were washed three times, resuspended in 1 ml of HBSS-BSA, and analyzed on a Coulter Epics Elite Cytometer. Non-specific binding of

the fluorescent secondary antibody was determined from samples lacking the primary antibody.

[199]        *Example III-5. Binding Assays to Intact Cells*

[200]        Assays with C441 cells were performed with cells grown to confluency in 24 well dishes. HUVEC cells were grown to confluency in 6 well dishes. Monolayers were incubated at 4°C for 2 hours with various amounts of mAb DC-101 in binding buffer (DMEM, 50 Mm HEPES pH 7.0, 0.5% bovine serum albumin). Cells were then washed with cold phosphate buffered saline (PBS) and incubated with a secondary anti-rat IgG antibody conjugated with biotin at a final concentration of 2.5 ug/ml. After 1 hour at 4°C cells were washed and incubated with a streptavidin-horse radish peroxidase complex for 30 minutes at 4°C. Following washing, cell-bound antibody was determined by measuring the absorbance at 540 nm obtained with a colormetric detection system (TMB, Kirkegaard and Perry). The OD 540 nm of the secondary antibody alone served as the control for non-specific binding.

[201]        *Example III-6. Cell proliferation assays*

[202]        Mitogenic assays were performed using the Cell Titer 96 Non Radioactive Cell Proliferation Assay Kit (Promega Corp., Madison, WI). In this assay proliferation is measured color metrically as the value obtained from the reduction of a tetrazolium salt by viable cells to a formazan product. Briefly, HUVEC cells were grown in 24 well gelatin-coated plates in EGM basal media at 1000 cells/well. After a 48-hour incubation various components were added to the wells. VEGF was added at 10 ng/ml to the media in the presence and absence of 1 µg/ml of mAb DC-101. Where indicated, heparin (Sigma) was added to a final concentration of 1 µg/ml. Cells were then incubated for an additional 3 days. To measure cell growth, a 20 µl aliquot of tetrazolum dye was added to each well and cells were incubated for 3 hrs at 37°C. Cells were solubilized and the absorbance (OD570) of the formazan product was measured as a quantitation of proliferation.

[203] *EXAMPLE IV. IN VITRO ACTIVITY ASSAYS*

[204] *Example IV-1. Murine anti-flk-1 (VEGFR-2) Mabs 25 and 73 elicit a specific neutralization of VEGF induced activation of the flk-1 (VEGFR-2)/fms receptor*

[205] Assays were performed with immunoprecipitated *FLK/fms* and PDGF receptors from equal concentrations of the flk-1 (VEGFR-2)/fms transfected 3T3 cell line, C441 whereas the human EGFR was immunoprecipitated from the tumor cell line, KB. Cells were stimulated with RPMI-0.5% BSA containing 20 ng/ml VEGF (flk-1/fms), DMEM-10% calf serum (PDGFR), or 10 ng/ml EGF (EGFR), in the presence and absence of 10 ug/ml of the murine anti-flk-1 Mabs, 25 and 73. Following stimulation, cells were washed with PBS-1mM sodium orthovanadate and lysed. Flk-1/fms and PDGFR were immunoprecipitated from lysates with peptide generated polyclonal antibodies against the C-terminal region of the c-fms (IM 133) and the PDGF (UBI) receptors, respectively. EGFR was immunoprecipitated with a Mab (C225) raised against the N-terminal region of the human receptor. Immunoprecipitated lysates were subjected to SDS polyacrylamide electrophoresis followed by western blotting. Blots were probed with an anti-PTyr Mab (UBI) to detect receptor activation. Receptor neutralization of stimulated cells was assessed relative to an irrelevant Mab and the unstimulated control.

[206] *Example IV-2. Detection of the flk-1 (VEGFR-2)/fms receptor by western blotting using Mab 25 and Mab 73 as probes*

[207] Receptor was detected by the murine anti-flk-1 (VEGFR-2) Mabs on western blots of the flk-1 (VEGFR-2)/fms receptor immunoprecipitated by a peptide generated polyclonal antibody against the C-terminal region of the c-fms receptor from lysate prepared from equal concentrations of transfected 3T3 cell line C441. Following analysis by SDS gel electrophoresis and western blotting, the blot was divided into four parts and each section was probed with 50 µg/ml of the anti-flk-1 (VEGFR-2) Mabs 25 and 73. Blots were then stripped and reprobed with the anti-fms polyclonal antibody to verify that the bands detected by each Mab represented the flk-1 (VEGFR-2)/fms receptor.

[208] *Example IV-3. Detection of activated KDR (VEGFR-2) from VEGF stimulated HUVEC and OVCAR-3 cells by immunoprecipitation with anti-flk-1 (VEGFR-2) Mabs*

[209] Proteins were immunoprecipitated by different antibodies from a lysate of freshly isolated HUVEC. Prior to lysis, cells were stimulated with 20 ng/ml VEGF for 10 minutes

at room temperature in RPMI-0.5% BSA and washed with PBS containing 1mM sodium orthovanadate. Individual immunoprecipitations were performed with equal volumes of lysate and then subjected to SDS polyacrylamide electrophoresis followed by western blotting. The blot was probed initially with an anti-PTyr Mab (UBI) and then sequentially stripped and reprobed with a peptide generated polyclonal antibody against the interkinase of flk-1/KDR (VEGFR-1/VEGFR-2, IM 142), followed by a polyclonal antibody against the C-terminal region of flk-1 (VEGFR-1) (Santa Cruz Biotechnology, Inc). The immunoprecipitations were performed with an irrelevant rat Mab, 23H7, an irrelevant mouse Mab, DAB 8, versus the anti-flk-1 (VEGFR-2) Mabs, DC-101, 73, 25 and an anti-flk-1/KDR (VEGFR-1/VEGFR-2) polyclonal antibody, IM 142. In some cases blots were stripped and reprobed with the anti-flk-1 Mabs 73 and 25 to detect cross reactive bands.

[210] A similar protocol was employed to detect KDR (VEGFR-2) receptor form(s) in the ovarian carcinoma cell line OVCAR-3.

[211] *EXAMPLE V. ACTIVITY OF ANTIBODIES*

[212] *Example V-1. ELISA and Immunoprecipitation with DC-101*

[213] Rat IgG1 monoclonal antibody DC-101 was found to be specific for the murine tyrosine kinase receptor flk-1 (VEGFR-2). ELISA data showed that the antibody bound to purified flk-1 (VEGFR-2):SEAP but not alkaline phosphatase or other receptor tyrosine kinases such as FLK-2. As seen in Figure 1, DC-101 immunoprecipitates murine flk-1 (VEGFR-2): SEAPS but not SEAPS alone.

[214] *Example V-2. Inhibition Of Flk-1 (VEGFR-2) Receptor Phosphorylation With DC-101*

[215] Experiments were then performed to determine whether DC-101 could neutralize phosphorylation of flk-1 (VEGFR-2) in C441 cells by its cognate ligand, VEGF<sub>165</sub>. In these studies, monoclonal antibody and VEGF were added simultaneously to monolayers for 15 minutes at room temperature. These conditions were designed to determine the competitive effects (competitive inhibition) of the antibody on receptor/ligand binding. The results of these assays, shown in Figure 2a, indicate that VEGF<sub>165</sub> induced phosphorylation of the flk-1 (VEGFR-2)/fms receptor was markedly reduced when cells were assayed in the presence of DC-101. In addition, these data suggest that the Mab

competes with VEGF<sub>165</sub> to prevent a full activation of receptor by ligand. To determine the sensitivity of the VEGF-flk-1 (VEGFR-2) interaction to inhibition by DC-101, C441 cells were assayed at maximal stimulatory concentrations of VEGF<sub>165</sub> (40 ng/ml) combined with varying levels of the antibody. The results of these Mab titrations are shown in Figure 2b. A marked decrease in the phosphorylation of flk-1 (VEGFR-2) by VEGF<sub>165</sub> was observed when DC-101 was included at concentrations greater than 0.5 µg/ml. These data show that relatively low concentrations of antibody (<1 µg/ml) are sufficient to inhibit receptor activation by ligand. At 5 µg/ml the antibody is able to neutralize VEGF<sub>165</sub> stimulation of flk-1 (VEGFR-2) in the presence of excess ligand at 80 ng/ml (Figure 3a and 3b). As a control, the effect of DC-101 was tested on the fully stimulated fms/*FLK-2* receptor (10A2 cell line) using CSF-1. Under these conditions, DC-101 showed no effect on receptor activation.

[216]        *Example V-3. Inhibition studies with DC-101*

[217]        The extent and specificity of Mab inhibition was further assessed by studies in which DC-101 was preincubated with cells before the addition of ligand to allow maximal interaction of antibody with receptor. In these experiments, monolayers were incubated with 5 µg/ml of DC-101, a rat anti-*FLK-2* Mab (2A13) prepared by conventional techniques (ImClone, NY), and control rat IgG1 (Zymed Labs) for 15 minutes at room temperature prior to the addition of 40 ng/ml of VEGF<sub>165</sub> for an additional 15 minutes. For comparison, assays were run in which DC-101 and VEGF<sub>165</sub> were added simultaneously (competitive inhibition). The results of these studies (Figure 4) show that preincubation of the anti-flk-1 (VEGFR-2) monoclonal antibody with flk-1 (VEGFR-2)/fms transfected cells completely abrogates receptor activation by VEGF<sub>165</sub>. Similar results were observed using VEGF<sub>121</sub> for stimulation. While phosphorylation of flk-1 (VEGFR-2) by VEGF is not affected by the addition of irrelevant isotype matched rat antibodies, the reactivity of the same blot probed with the anti-fms polyclonal antibody shows an equal level of receptor protein per lane. These data indicate that the inhibition of phosphorylation observed with DC-101 was due to the blockage of receptor activation rather than a lack of receptor protein in the test samples.

[218]        *Example V-4. Binding of DC-101 to C441 cells by FACS analysis*

[219]        The mAb was assayed by FACS analysis for binding to 3T3 cells transfected with the flk-1 (VEGFR-2)/fms receptor (C441 cells). The results, shown in Figure 6, demonstrate that the chimeric flk-1 (VEGFR-2)/fms expressed on the surface of C441 cells is specifically recognized by mAb DC-101 and not by an antibody of the same isotype raised against the related tyrosine kinase receptor, *FLK-2*. The efficacy of the mAb-receptor interaction at the cell surface was determined from assays in which varying levels of mAb binding was measured on intact C441 cells. These results, shown in Figure 7, indicate that mAb binds to the flk-1 (VEGFR-2)/fms receptor with a relative apparent affinity of approximately 500 ng/ml. These results indicate that the mAb has a strong affinity for cell surface expressed flk-1 (VEGFR-2).

[220]        *Example V-5. Reactivity of DC-101 by Immunoprecipitation*

[221]        The extent of DC-101 reactivity with the flk-1 (VEGFR-2)/fms receptor was further assessed by determining the capacity of the antibody to immunoprecipitate the receptor following activation by VEGF. Figure 8 shows an immunoprecipitation by mAb DC-101 of the phosphorylated flk-1 (VEGFR-2)/fms receptor from VEGF stimulated C441 cells. The results show that the DC-101 monoclonal and anti-fms polyclonal antibodies display similar levels of receptor interaction while rat anti *FLK-2* antibodies 2H37 (IgG1) and 2A13 (IgG2a) show no reactivity. Experiments were then performed to determine whether mAb DC-101 could neutralize the VEGF induced phosphorylation of flk-1 (VEGFR-2)/fms at maximal stimulatory concentrations of ligand (40 ng/ml). In these studies, monoclonal antibody was added to monolayers either simultaneously with ligand or prior to ligand stimulation and assayed for 15 minutes at room temperature. These conditions were studied to determine both the competitive effects (competitive inhibition) of the antibody on receptor/ligand binding as well as the efficacy of prebound antibody to prevent receptor activation. The results of these assays, shown in Figure 4, indicate that phosphorylation of the flk-1 (VEGFR-2)/fms is reduced by the simultaneous addition of mAb with VEGF and markedly inhibited by antibody prebound to the receptor. A densitometry scan of these data revealed that mAb DC-101 interacts with flk-1 (VEGFR-2)/fms to inhibit phosphorylation to a level that is 6% (lane 5, P) and 40% (lane 6, C) of the fully stimulated receptor control (lane 4). From these data we infer that mAb

DC-101 strongly competes with the ligand-receptor interaction to neutralize flk-1 receptor activation. To determine the sensitivity of the VEGF-flk-1 (VEGFR-2) interaction to inhibition by mAb DC-101, C441 cells were assayed with maximal VEGF levels in the presence of increasing concentrations of antibody. Assays were performed with the mAb under competitive and prebinding conditions. The results of these mAb titrations are shown in Figure 9. A marked decrease in the phosphorylation of flk-1 (VEGFR-2) is observed when mAb DC-101 competes with VEGF antibody at concentrations greater than 0.5  $\mu\text{g/ml}$ . These data also show that relatively low concentrations of prebound antibody ( $<1 \mu\text{g/ml}$ ) are sufficient to completely inhibit receptor activation by ligand.

[222]        *Example V-6. Activity of DC-101 by phosphorylation assay*

[223]        To further evaluate the antagonistic behavior of mAb DC-101 on receptor activation, phosphorylation assays were performed in which a fixed amount of antibody (5  $\mu\text{g/ml}$ ) was added to C441 cells stimulated with increasing amounts of ligand (Figure 3a). The level of phosphorylation induced by each ligand concentration in the presence and absence of mAb DC-101 was also quantitated by densitometry readings. The plot of these data given in Figure 3b indicates that the antibody was able to partially neutralize receptor phosphorylation even in the presence of excess amounts of VEGF. To evaluate the specificity of mAb DC-101 on receptor activation, the antibody was tested for its ability to competitively inhibit CSF-1 induced activation of the *fms/FLK-2* receptor in the 3T3 transfected cell line, 10A2. In these experiments 5  $\mu\text{g/ml}$  of mAb DC-101 was tested together with CSF-1 concentrations (20-40  $\text{ng/ml}$ ) that are known to result in full activation of the receptor. These results, which are shown in Figure 10, indicate that mAb DC-101 has no effect on the CSF-1 mediated phosphorylation of the *fms/FLK-2* receptor.

[224]        *Example V-7. DC-101 inhibition by pre-incubation studies*

[225]        The extent and specificity of antibody inhibition was further assessed by studies in which DC-101 or an irrelevant antibodies were preincubated with cells before the addition of ligand to assure maximal interaction of antibody with receptor. In these experiments, monolayers were preincubated with either 5  $\mu\text{g/ml}$  of DC-101, a rat anti-*FLK-2* mAb (2A13) or a control rat IgG1 (Zymed Labs) prior to the addition of 40  $\text{ng/ml}$  of VEGF. For comparison, competitive assays were run in which antibodies and VEGF were added simultaneously. The results of these studies show that only the preincubation of the anti-

flk-1 (VEGFR-2) monoclonal antibody with flk-1 (VEGFR-2)/fms transfected cells completely abrogates receptor activation by VEGF while phosphorylation of flk-1 (VEGFR-2) by VEGF is not affected by the addition of irrelevant isotype matched rat antibodies. The reactivity of the same blot probed with the anti-fms polyclonal (Figure 11) shows an equal level of receptor protein per lane. These data indicate that the lack of phosphorylation observed with mAb DC-101 treated cells was due to the blockage of a VEGF-induced phosphorylation of equal amounts of expressed receptor.

[226]           *Example V-8. Interaction of antibodies with homologous receptor forms*

[227]           Experiments were then conducted to determine whether the anti-flk-1 (VEGFR-2) monoclonal antibodies interact with homologous receptor forms on human endothelial cells. A titration of increasing concentrations of DC-101 on cloned HUVEC cells (ATCC) indicated that the antibody displayed a complex binding behavior. The data represent differential antibody interactions with VEGF receptors reported to occur on endothelial cells (Vaisman et al., J. Biol. Chem. 265, 19461-19466, 1990). The specificity of DC-101 interaction with VEGF stimulated HUVEC cells was then addressed using phosphorylation assays under similar conditions as those reported for Figure 8. In these studies DC-101 immunoprecipitates protein bands from HUVEC cells that have molecular weights similar to those reported for cross linked VEGF-receptor bands when the ligand component is subtracted (Figure 12). These bands display an increased phosphorylation when cells are stimulated by VEGF (compare lanes 1 and 2 in Figure 12). In addition, the VEGF induced phosphorylation of the receptor bands is potentiated by the inclusion of 1 µg/ml heparin in the assay (lane 3 in Figure 12). These findings are consistent with previous reports of increased VEGF binding to endothelial cells in the presence of low concentrations of heparin (Gitay-Goren et al., J. Biol. Chem. 267, 6093-6098, 1992).

[228]           It is difficult to ascertain which immunoprecipitated protein interacts with DC-101 to generate the complex of phosphorylated bands observed in Figure 12 given the various receptor forms shown to bind VEGF on HUVEC and the possibility of their association upon stimulation. Cell surface expressed receptor forms with molecular weights of approximately 180 (KDR (VEGFR-2)), 155, 130-135, 120-125 and 85 have been reported to bind VEGF on HUVEC. Such findings address the possibility that several different receptor forms may heterodimerize upon ligand stimulation in a manner similar to that

reported for KDR-flk-1 (VEGFR-2/VEGFR-1). However, with the exception of KDR (VEGFR-2), the exact nature and role of these receptor forms have yet to be defined. Consequently, antibody reactivity may result from interaction(s) with one of several VEGF receptors independent of KDR (VEGFR-2).

[229] DC-101 does not react with human KDR (VEGFR-2) in an ELISA format nor bind to freshly isolated HUVEC by FACS analysis. These results suggest that a direct interaction of DC-101 with human KDR (VEGFR-2) is highly unlikely.

[230] Unlike DC-101, Mab 25 and Mab 73 both react with human KDR (VEGFR-2) in an ELISA format and bind to freshly isolated HUVEC by FACS analysis.

[231] *Example V-9. Mitogenic assays of HUVEC.*

[232] An inhibitory effect of DC-101 on endothelial cells was observed when the antibody was tested in mitogenic assays of HUVEC cells (ATCC) stimulated with VEGF in the presence and absence of antibody (Figure 12). These results show that a marked increase in cell proliferation by VEGF is reduced approximately 35% by DC-101. Heparin shows no differential effect on cell growth under the growth conditions employed in these assays.

[233] Since DC-101 can exert effects on VEGF induced proliferation and receptor phosphorylation of HUVEC it is conceivable that these results are due to a Mab interaction with an undefined receptor form which is poorly accessible at the cell surface, but which plays some role, albeit minor, in HUVEC growth. Also, the immunoprecipitation of phosphorylated bands of the correct molecular weight by DC-101 from VEGF stimulated HUVEC also supports the notion that DC-101 may interact with an undefined flk-1 (VEGFR-2) like protein that associates with an activated receptor complex.

[234] *Example V-10. Binding of Mab 25 and Mab 73 to C441 cells and HUVEC*

[235] Mabs 25 and 73 bind to C441 and HUVEC by FACS analysis and show internalization in both cell lines. Results from western blots show that both anti-flk-1 Mabs can detect the band(s) for the FLK/fms receptor in immunoprecipitates by an anti-fms polyclonal antibody from C441 cells. (See example IV-2 above for protocol.) These

antibodies elicit a specific neutralization of VEGF induced activation of the flk-1 (VEGFR-2)/fms receptor and have no effect on the phosphorylation of the mouse PDGF receptor by PDGF or the human EGF receptor by EGF. (See example IV-1 above for protocol.) They have the capacity to inhibit VEGF stimulated HUVEC in proliferation assays to 50% whereas DC-101 affects growth to a far lesser extent.

[236] *Example V-11. Immunoprecipitation of KDR (VEGFR-2) with Mab25 and Mab73*

[237] KDR (VEGFR-2) represents one of the phosphoproteins immunoprecipitated by the Mab25 and Mab 73 from activated HUVEC. KDR (VEGFR-2) was detected in western blot and immunoprecipitation analyses using an anti-flk-1/KDR (VEGFR-1/VEGFR-2) polyclonal antibody (IM142) from VEGF-stimulated early passage HUVEC. Conversely, bands immunoprecipitated by these antibodies from VEGF-stimulated HUVEC are cross reactive with IM142 but not an anti-flt-1 (anti-VEGFR-1) polyclonal antibody. These findings infer that the Mabs may affect the activity of KDR (VEGFR-2) in HUVEC based on experimental evidence implicating KDR (VEGFR-2) as the VEGF receptor responsible for the proliferative response in activated endothelial cells. (See example IV-3 above for protocol.)

[238] *EXAMPLE VI. PRESENCE OF VEGF RECEPTOR FORMS ON NON-ENDOTHELIAL (TUMOR) CELLS*

[239] Several tumor lines were screened for protein reactivity with DC-101 by immunoprecipitation and detection with antiphosphotyrosine. Immunoblots from the cell lines 8161 (melanoma) and A431 (epidermoid carcinoma) yielded phosphorylated bands with molecular weights of approximately 170 and 120 kD. These results indicate that a human VEGF receptor form is expressed in non-endothelial cells, such as tumor cells.

[240] Similar experiments have shown that a KDR (VEGFR-2) like receptor is expressed in an ovarian carcinoma cell line, OVCAR-3. These cells also appear to secrete VEGF. Phosphorylated bands are immunoprecipitated by an anti-KDR (VEGFR-2) polyclonal antibody from VEGF-stimulated OVCAR-3 cells that are reactive with anti-flk-1 (VEGFR-2) Mabs by western blotting. Also, bands immunoprecipitated by the murine Mabs from these cells show cross reactivity with the same polyclonal antibody. Furthermore certain murine anti-flk-1 (VEGFR-2) Mabs elicit an inhibitory effect on these cells in proliferation assays. These results demonstrate nonendothelial expression (i.e. on

tumor cells) of human VEGF receptor forms. The data from the phosphorylation and proliferation assays also suggest that VEGF can modulate receptor activity in an autocrine and paracrine manner during tumorigenesis. (See Example IV-3 above for protocol.)

[241] *EXAMPLE VII. IN VIVO STUDIES USING DC-101*

[242] *Example VII-1. inhibition in vivo of angiogenesis by DC-101*

[243] *In vivo* studies were designed to determine if an anti-flk-1 (VEGFR-2) monoclonal antibody would block the growth of VEGF-expressing tumor cells. In these experiments, a human glioblastoma multiform cell line was used that has high levels of VEGF message and secretes about 5 ng/ml of VEGF growth factor after a 24 hour conditioning in serum free medium (Figure 5).

[244] On day zero, athymic nude mice (nu/nu; Charles River Labs) were injected in the flank with 1-2 million glioblastoma cells. Beginning on the same day, animals received intraperitoneal injections of either DC-101 and control antibodies (100 ug/animal). The mice received subsequent antibody treatments on days 3, 5, 7, 10, 12, 14, 17, 19, and 21. Animals received injections of 100 µg of either DC-101 or a control rat antibody to the murine *FLK-2* (2A13) receptor on days 0, 3, 5, 7, 10, 12, 14, 17, 19, and 21 for a total inoculation of 1 mg/animal. Tumors began to appear by day 5 and followed for 50 days. Tumor size was measured daily with a caliper and tumor volume calculated by the following formula:  $\frac{p}{6} \times \text{larger diameter} \times (\text{smaller diameter})^2$  (Baselga J. Nat'l Cancer Inst. 85: 1327-1333). Measurements were taken at least three times per week and tumor volume calculated as described above. One tumor bearing animal in the DC-101 group died early in the study and was not used to determine statistical significance between the groups.

[245] Figures 14a and 14b show a comparison between the DC-101 and the control 2A13 group of reduction in tumor growth over 38 days in individual animals. Although all animals developed tumors of varying sizes and number during the course of the study, DC-101-treated mice showed an overall delay in tumor progression. One mouse in the DC-101 group remained tumor free until day 49 when a small growth was observed. Even then, tumor growth was markedly suppressed. Statistical analysis of the data was done to assess differences in tumor size between the two groups. Data was subjected to a standard

analysis of covariance where tumor size was regressed on time with treatment as a covariate. The results showed that reduction in tumor size over time for the DC-101 group was significantly different ( $p < 0.0001$ ) from that seen for 2A13 injected mice.

[246] Figure 15 shows the therapeutic efficacy of DC-101 in athymic nude mice transplanted with the human glioblastoma tumor cell line GBM-18, which secretes VEGF. Nude mice were injected subcutaneously with GBM-18 cells and divided into three groups of treatment: a PBS control, an irrelevant rat IgG1 control, and DC-101. Treatments were administered simultaneously with tumor xenografts and continued for four weeks. The results showed that GBM-18 tumor growth in DC-101-treated nude mice was significantly reduced relative to controls. This experiment indicates that DC-101 suppresses tumor growth by blocking VEGF activation of flk-1 (VEGFR-2) on tumor associated vascular endothelial cells, and that DC-101 has therapeutic value as an anti-angiogenic reagent against vascularized tumors secreting VEGF.

[247] Monoclonal antibodies to flk-1 (VEGFR-2) receptor tyrosine kinase inhibit tumor invasion by abrogating angiogenesis. Invasive growth and angiogenesis are essential characteristics of malignant tumors. Both phenomena proved to be suitable to discriminate benign from malignant keratinocytes in a surface transplantation assay. After transplantation of a cell monolayer attached to a collagen gel onto the back muscle of nude mice, all tumor cells initially formed organized squamous epithelia, but only malignant keratinocytes grew invasively within 2-3 weeks. Both benign and malignant cells induced angiogenesis. Angiogenic response to malignant cells, however, occurred earlier, is much stronger, and capillary growth directed toward malignant epithelia. Moreover, in transplants of benign tumor cells, capillaries regressed after 2-3 weeks, whereas malignant keratinocytes maintain the level of ongoing angiogenesis. The vascular endothelial growth factor (VEGF) and its cognate receptor play a pivotal role in tumor angiogenesis. The administration of DC-101 disrupted ongoing angiogenesis leading to inhibition of tumor invasion. The antibody prevented maturation and further expansion of newly formed vascular network, but did not significantly interfere with initial angiogenesis induction. These results provide evidence that tumor invasion requires precedent angiogenesis, and that the VEGF receptors are crucial in maintaining angiogenesis in this model system.

[248] *Example VII-2. Effect of different concentrations of DC-101 on established glioblastoma (gbm-18) tumors*

[249] Athymic mice (nu/nu) were inoculated subcutaneously with GBM-18 (human glioblastoma multiformae). Antibody therapy was initiated when the tumors reached an average volume of 100-200 mm<sup>3</sup>. Treatment consisted of six injections (twice weekly for 3 weeks) of the following: (i) DC-101 at 200, 400 or 800 µg/injection; (ii) an irrelevant isotype matched rat IgG (400 µg/injection); or, (iii) PBS. Tumor volumes were measured with a caliper. Tumor inhibition in the DC-101 groups was found to be significant (\*) vs. the PBS and irrelevant monoclonal antibody groups.

[250] Another experiment demonstrates the effects of the rat anti-flk-1 (VEGFR-2) monoclonal antibody DC-101 on the growth of GBM-18 tumors in nude mice. Animals (nu/nu; Charles River Labs; ten animals per group) were injected subcutaneously with GBM-18 cells (human glioblastoma [100]; 1 million per animal) on day 0. Treatments with PBS or DC-101 (200 µg per injection) were begun on day 7 and continued twice weekly for 3 weeks (6x). Graphs show a plot of the mean tumor volumes and regressed data for each group over time with their respective tumor growth rates (slopes given as  $\lambda$ ; solid lines) and 99% confidence limits (dotted lines). The slope of the line for animals treated with DC-101 was significantly different from that of PBS ( $p < 0.01$ ). It is important to note that an irrelevant rat IgG1 monoclonal antibody (anti-mouse IgA; Pharmigen) had no effect on the growth of GBM-18 xenografts and gave results similar to that observed with PBS (data not shown).

[251] *EXAMPLE VIII. ANTI-flk-1 (VEGFR-2) ANTIBODY SELECTIVELY INCREASES RADIATION-INDUCED CURE RATE OF HUMAN TUMOR XENOGRAFTS IN NUDE MICE*

[252] This example evaluates whether the monoclonal antibody DC-101 blocking the crucial VEGF receptor-2, flk-1 (VEGFR-2), on murine endothelial cells of tumor vessels increases curability of tumor xenografts by fractionated radiotherapy (RT), and whether the antibody concurrently modulates the radiation reaction of normal tissue (mouse skin).

[253] **Materials & Methods:** The human small cell lung carcinoma 54A and glioblastoma multiforme U87 were implanted subcutaneously into the hind leg of nude mice. Treatment was begun when a tumor reached 8mm in diameter (day 0). DC-101 was injected intraperitoneally every 3 days at a dose of 20 or 40 mg/kg body-weight, for a total

of 6 injections. Graded total doses of radiation were given in equal daily fractions on 5 consecutive days. On day 0, a mouse received the first injection of the antibody, or RT was started. For the combined treatment, DC-101 administration was commenced on day 0, and RT was begun on day 1. Tumor size was measured 2-3 times a week after treatment. The mice with locally controlled tumors were followed-up for 90 days after the last tumor recurrence was observed in any group. Acute reaction of skin in the field of tumor irradiation was evaluated using a scoring scale during the first 30 days after the beginning of RT.

[254]        Results: The antibody used alone induced growth inhibition (but not regression) of both tumors in a dose-dependent manner. The effect was more pronounced in 54A than in U87 xenografts. In combination with the lowest doses of radiation (25-30 Gy total), DC-101 provided an additional tumor growth delay when compared with RT alone, in either model. The antibody, also in a dose-dependent fashion, augmented the curative effect of RT. For example, at its higher dose, DC-101 decreased the dose of radiation necessary to control 50% of tumors locally: 1.7 fold in 54A xenografts (from 67.6 Gy for RT alone to 39.1 Gy for the combined therapy), and 1.3 fold in U87 (from 97.8 to 74.8 Gy). It is also of particular importance that such effects of DC-101 were selective for tumors. That is, no parallel changes of skin radiation reaction by the antibody were detected. As assessed in additional experiments, the DC-101-induced enhancement of the radiation response of tumors was not associated with their radiosensitization or changes in oxygenation, while correlated with a significant decrease of the tumor interstitial fluid pressure by the antibody.

[255]        Conclusion: The results collectively suggest that the blockage of VEGF-signaling pathways by an antibody against the main receptor to these growth factor molecules can selectively potentiate the tumor curative response to fractionated RT; and thus, provide a therapeutic gain.

[256] *EXAMPLE IX. PRODUCING SINGLE CHAIN ANTIBODIES*

[257] *Example IX-1 Materials*

[258] *Example IX-1 (a). Cell lines and Proteins*

[259] Primary-cultured HUVEC was maintained in EBM-2 medium at 37°C, 5% CO<sub>2</sub>. Cells were used between passage 2-5 for all assays. VEGF<sub>165</sub> protein was expressed in baculovirus and purified. Complementary DNA encoding the extracellular domain of KDR (VEGFR-2) was isolated by RT-PCR from human fetal kidney mRNA and subcloned into the *Bgl* II and *Bsp*E I sites of the vector AP-Tag. In this plasmid the cDNA for KDR (VEGFR-2) extracellular domain is fused in-frame with the cDNA for human placental AP. The plasmid was electroporated into NIH 3T3 cells together with the neomycin expression vector pSV-Neo and stable cell clones were selected with G418. The soluble fusion protein KDR-AP was purified from cell culture supernatant by affinity chromatography using immobilized monoclonal antibodies to AP.

[260] *Example IX-1 (b). Mice immunization and construction of single chain antibody phage display library*

[261] Female BALB/C mice were given two intraperitoneal (i.p.) injections of 10 µg KDR-AP in 200 µl of RIBI Adjuvant System followed by one i.p. injection without RIBI adjuvant over a period of two months. The mice were also given a subcutaneous (s.c.) injection of 10 µg KDR-AP in 200 µl of RIBI at the time of the first immunization. The mice were boosted i.p. with 20 µg of KDR-AP three days before euthanasia. Spleens from donor mice were removed and the cells were isolated. RNA was extracted and mRNA was purified from total RNA of splenocytes. A scFv phage display library was constructed using the mRNA which was displayed on the surface of the filamentous phage M13.

[262] In displaying the scFv on filamentous phage surface, antibody V<sub>H</sub> and V<sub>L</sub> domains are joined together by a 15 amino-acid-long linker (GGGGS)<sup>3</sup> and fused to the N-terminal of phage protein III. A 15 amino-acid-long E tag, which is followed by an amber codon (TAG), was inserted between the C-terminal of V<sub>L</sub> and the protein III for detection and other analytic purposes. The amber codon positioned between the E tag and the protein III enables the construct to make scFv in surface-displaying format when transformed into a

suppressor host (such as TGI cells) and scFv in soluble form when transformed into a nonsuppressor host (such as HB2151 cells).

[263] The assembled scFv DNA was ligated into the pCANTAB 5E vector. The transformed TG1 cells were plated onto 2YTAG plates and incubated. The colonies were scraped into 10 ml of 2YT medium, mixed with 5 ml 50% glycerol and stored at -70°C as the library stock.

[264] *Example IX -1(c). Biopanning*

[265] The library stock was grown to log phase, rescued with M13K07 helper phage and amplified overnight in 2YTAK medium (2YT containing 100 µg/ml of ampicillin and 50 µg/ml of kanamycin) at 30°C. The phage preparation was precipitated in 4% PEG/0.5M NaCl, resuspended in 3% fat-free milk/PBS containing 500 µg/ml of AP protein and incubated at 37°C for 1 h to capture phage displaying anti-AP scFv and to block other nonspecific binding.

[266] KDR-AP (10 µg/ml) coated Maxisorp Star tubes (Nunc, Denmark) were first blocked with 3% milk/PBS at 37°C for 1 h, and then incubated with the phage preparation at room temperature for 1 h. The tubes were washed 10 times with PBST followed by 10 times with PBS (PBS containing 0.1% Tween 20). The bound phage was eluted at room temperature for 10 min. with 1 ml of a freshly prepared solution of 100 mM triethylamine. The eluted phage were incubated with 10 ml of mid-log phase TG1 cells at 37°C for 30 min. stationary and 30 min. shaking. The infected TG1 cells were then plated onto 2YTAG plates and incubated overnight at 30°C.

[267] Ninety-nine percent (185/186) of clones screened after the third round of panning were found to be specific KDR (VEGFR-2) binders. However, only 15 (8%) of these binders could block KDR (VEGFR-2) binding to immobilized VEGF. DNA BstNI fingerprinting of these 15 clones indicated the presence of 2 different digestion patterns; whereas 21 randomly picked VEGF nonblockers yielded 4 different patterns. All the digestion patterns were also seen in clones identified after the second round of panning. Representative clones of each digestion pattern were picked from clones recovered after the 2nd round of panning and subject to DNA sequencing. Out of 15 clones sequenced, 2 unique VEGF blockers and 3 nonblockers were identified. One scFv, p2A7, which neither

binds to KDR (VEGFR-2) nor blocks VEGF binding to KDR (VEGFR-2), was selected as a negative control for all studies.

[268]                    *Example IX-1 (d). Phage ELISA*

[269]                    Individual TG1 clones were grown at 37°C in 96 well plates and rescued with M13K07 helper phage as described above. The amplified phage preparation was blocked with 1/6 volume of 18% milk/PBS at RT for 1 h and added to Maxi-sorp 96-well microtiter plates (Nunc) coated with KDR-AP or AP (1 µg/ml x 100 µl). After incubation at room temperature for 1 h, the plates were washed 3 times with PBST and incubated with a rabbit anti-M13 phage Ab-HRP conjugate. The plates were washed 5 times, TMB peroxidase substrate added, and the OD at 450 nm read using a microplate reader and scFv antibodies were identified and sequenced.

[270]                    *Example IX-1(e). Preparation of soluble scFv*

[271]                    Phage of individual clones were used to infect a nonsuppressor *E.coli* host HB2151 and the infectant selected on 2YTAG-N plates. Expression of scFv in HB2 151 cells was induced by culturing the cells in 2YTA medium containing 1 mM isopropyl-1-thio-B-D-galactopyranoside at 30°C. A periplasmic extract of the cells was prepared by resuspending the cell pellet in 25 mM Tris (pH 7.5) containing 20% (w/v) sucrose, 200 mM NaCl, 1 mM EDTA and 0.1 mM PMSF, followed by incubation at 4°C with gentle shaking for 1 h. After centrifugation at 15,000 rpm for 15 min., the soluble scFv was purified from the supernatant by affinity chromatography using the RPAS Purification Module (Pharmacia Biotech).

[272]                    *Example IX-2. Assays*

[273]                    *Example IX-2 (a). Quantitative KDR (VEGFR-2) binding assay*

[274]                    Two assays were employed to examine quantitatively the binding of purified soluble scFv to KDR (VEGFR-2).

[275]                    Four different clones, including the two VEGF blockers, pIC11 and pIF12, one nonblocker, the dominant clone p2A6 and the nonbinder p2A7, were expressed in shaker flasks using a nonsuppressor host *E.coli* HB2151 cells. The soluble scFv was purified

from the periplasmic extracts of *E.coli* by anti-E-tag affinity chromatography. The yield of purified scFv of these clones ranged from 100 - 400 µg / liter culture.

[276] In the direct binding assay, various amounts of soluble scFv were added to KDR (VEGFR-2)-coated 96-well Maxi-sorp microtiter plates and incubated at room temperature for 1 h, after which the plates were washed 3 times with PBST. The plates were then incubated at room temperature for 1 h with 100 µl of mouse anti-E tag antibody followed by incubation with 100 p1 of rabbit anti-mouse antibody-HRP conjugate. The plates were washed and developed following the procedure described above for the phage ELISA.

[277] In another assay, i.e., the competitive VEGF blocking assay, various amounts of soluble scFv were mixed with a fixed amount of KDR-AP (50 ng) and incubated at room temperature for 1 h. The mixture were then transferred to 96-well microtiter plates coated with VEGF<sub>165</sub> (200 ng/well) and incubated at room temperature for an additional 2 h, after which the plates were washed 5 times and the substrate for AP was added to quantify the bound KDR-AP molecules. IC<sub>50</sub>, i.e., the scFv concentration required for 50% inhibition of KDR (VEGFR-2) binding to VEGF, was then calculated.

[278] Figure 16 shows the dose-dependent binding of scFv to immobilized KDR (VEGFR-2) as assayed by a direct binding ELISA. Clone p1C11 and p1F12, but not p2A6, also block KDR (VEGFR-2) binding to immobilized VEGF as shown in Fig. 17. Data shown in Figure 17 are the means ± SD of triplicate determinations. The negative control clone, p2A7, did not bind to KDR (VEGFR-2) nor block KDR (VEGFR-2) binding to VEGF (Fig. 16 and 17). Clone p1C11, the dominant clone after each round of panning, showed the highest KDR (VEGFR-2) binding capacity and the highest potency in blocking VEGF binding to KDR (VEGFR-2) (Table 1). The antibody concentrations of clone p1C11 required for 50% of maximum binding to KDR (VEGFR-2) and for 50% of inhibition of KDR (VEGFR-2) binding to VEGF (Fig 17) were 0.3 nM and 3 nM, respectively (See Table 1). FACS analysis demonstrated that p1C11, p1F12 and p2A6 were also able to bind to cell surface expressed receptor on HUVEC.

[279] *Example IX-2 (b). BIAcore analysis of the soluble scFv*

[280] The binding kinetics of soluble scFv to KDR (VEGFR-2) were measured using BIAcore biosensor (Pharmacia Biosensor). KDR-AP fusion protein was immobilized onto

a sensor chip and soluble scFv were injected at concentrations ranging from 62.5 nM to 1000 nM. Sensorgrams were obtained at each concentration and were evaluated using a program, BIA Evaluation 2.0, to determine the rate constant *kon* and *koff*. *Kd* was calculated from the ratio of rate constants *koff/kon*.

[281] Table 1 shows the results of the surface plasmon resonance on a BIAcore instrument. The VEGF-blocking scFv, p1C11 and p1F12, bound to immobilized KDR (VEGFR-2) with *Kd* of 2.1 and 5.9 nM, respectively. The non-blocking scFv, p2A6, bound to KDR (VEGFR-2) with approximately a 6-fold weaker affinity (*Kd*, 11.2 nM) than the best binder p1C11, mainly due to a much faster dissociation rate. As anticipated, p2A7 did not bind to the immobilized KDR (VEGFR-2) on the BIAcore.

[282] *Example IX-2 (c). Phosphorylation assay*

[283] Phosphorylation assays were performed with early passage HUVEC following a protocol described previously. Briefly, HUVEC were incubated in serum free EBM-2 base medium supplemented with 0.5% bovine serum albumin at room temperature for 10 min. in the presence or absence of scFv antibodies at 5 µg/ml, followed by stimulation with 20 ng/ml VEGF<sub>165</sub> at room temperature for an additional 15 min. The cells were lysed and the KDR (VEGFR-2) receptor was immunoprecipitated from the cell lysates with Protein A Sepharose beads coupled to a rabbit anti-KDR (anti-VEGFR-2) polyclonal antibody (ImClone Systems Incorporated). The beads were washed, mixed with SDS loading buffer, and the supernatant subjected to Western blot analysis. To detect KDR (VEGFR-2) phosphorylation, blots were probed with an anti-phosphotyrosine Mab, 4G10. For the MAP kinase activity assay, cell lysates were resolved with SDS-PAGE followed by Western blot analysis using a phospho-specific MAP kinase antibody. All signals were detected using ECL.

[284] Results showed that VEGF-blocking scFv p1C11, but not the non-blocking scFv p2A6, was able to inhibit KDR (VEGFR-2) receptor phosphorylation stimulated by VEGF. Further, p1C11 also effectively inhibited VEGF-stimulated activation of MAP kinases p44/p42. In contrast, neither p1C11, nor p2A6 inhibited FGF-stimulated activation of MAP kinases p44/p42.

[285] *Example IX-2 (d). Anti-mitogenic assay.*

[286] HUVEC (5 x 10<sup>3</sup> cells/well) were plated onto 96-well tissue culture plates (Wallach, Inc., Gaithersburg, MD) in 200  $\mu$ l of EBM-2 medium without VEGF, bFGF or EGF and incubated at 37°C for 72 h. Various amounts of antibodies were added to duplicate wells and pre-incubated at 37°C for 1 h, after which VEGF<sub>165</sub> was added to a final concentration of 16 ng/ml. After 18 h of incubation, 0.25  $\mu$ Ci of [<sup>3</sup>H]-TdR (Amersham) was added to each well and incubated for an additional 4 h. The cells were placed on ice, washed twice with serum-containing medium, followed by a 10 minute incubation at 4°C with 10% TCA. The cells were then washed once with water and solubilized in 25  $\mu$ l of 2% SDS. Scintillation fluid (150  $\mu$ l/well) was added and DNA incorporated radioactivity was determined on a scintillation counter (Wallach, Model 1450 Microbeta Scintillation Counter).

[287] The ability of scFv antibodies to block VEGF-stimulated mitogenic activity on HUVEC is shown in Fig. 18. The VEGF-blocking scFv p1C11 strongly inhibited VEGF induced DNA synthesis in HUVEC with an EC<sub>50</sub>, i.e., the antibody concentration that inhibited 50% of VEGF-stimulated mitogenesis of HUVEC, of approximately 5 nM. The non-blocking scFv p2A6 showed no inhibitory effect on the mitogenic activity of VEGF. Neither p1C11 nor p2A6 inhibited bFGF-induced DNA synthesis in HUVEC (not shown). Data shown in Fig. 18 are representative of at least three separate experiments. (!) VEGF only; (  $\nabla$  ) no VEGF.

[288] *Example IX-3. Producing Chimeric Antibodies from p1C11*

[289] *Example IX-3(a). Cell lines and Proteins*

[290] Primary-cultured human umbilical vein endothelial cells (HUVEC) were maintained in EBM-2 medium at 37°C, 5% CO<sub>2</sub>. Cells between passage 2-5 were used for all assays. VEGF<sub>165</sub> and KDR (VEGFR-2)-alkaline phosphatase fusion proteins (KDR-AP) were expressed in baculovirus and NIH 3T3 cells, respectively, and purified following the procedures described above. The anti-KDR (anti-VEGFR-2) scFv p1C11 and scFv p2A6, an antibody that binds to KDR (VEGFR-2) but does not block KDR (VEGFR-2)-VEGF interaction, were isolated from a phage display library constructed

from a mouse immunized with KDR (VEGFR-2) as described above. C225 is a chimeric IgG1 antibody directed against epidermal growth factor (EGF) receptor. See above.

[291] *Example IX-3(b). Cloning of the variable domains of scFv p1C11*

[292] The variable domains of the light ( $V_L$ ) (SEQ ID NO: 8 and SEQ ID NO: 16) and the heavy ( $V_H$ ) (SEQ ID NO: 7 and SEQ ID NO: 15) chains of p1C11 were cloned from the scFv expression vector by PCR using primers 1 and 2, and primers 3 and 4, respectively. The leader peptide sequence for protein secretion in mammalian cells was then added to the 5' of the  $V_L$  and the  $V_H$  by PCR using primers 5 and 2, and primers 5 and 4, respectively.

[293] Primer 1: 5' CTA GTA GCA ACT GCA ACT GGA GTA CAT TCA GAC ATC GAG CTC3' [SEQ ID No: 37]

[294] Primer 2: 5' TCG ATC TAG AAG GAT CCA CTC ACG TTT TAT TTC CAG3' *Bam*HI [SEQ ID NO: 38]

[295] Primer 3: 5' CTA GTA GCA ACT GCA ACT GGA GTA CAT TCA CAG GTC AAG CTG3' [SEQ ID No: 39]

[296] Primer 4: 5' TCG AAG GAT CCA CTC ACC TGA GGA GAC GGT3' *Bam*HI [SEQ ID No: 40]

[297] Primer 5: 5' GGT CAA AAG CTT ATG GGA TGG TCA TGT ATC ATC CTT TTT *Hind* III CTA GTA GCA ACT3' [SEQ ID No: 41]

[298] *Example IX-3(c). Construction of the expression vectors for the chimeric p1C11 IgG.*

[299] Separate vectors for expression of chimeric IgG light chain and heavy chains were constructed. The cloned  $V_L$  gene was digested with *Hind* III and *Bam*HI and ligated into the vector pKN100 containing the human  $\kappa$  light chain constant region ( $C_L$ ) to create the expression vector for the chimeric p1C11 light chain, c-p1C11-L. The cloned  $V_H$  gene was digested with *Hind* III and *Bam*HI and ligated into the vector pGID105 containing the human IgG1 ( $\gamma$ ) heavy chain constant domain ( $C_H$ ) to create the expression vector for

the chimeric p1C11 heavy chain, c-p1C11-H. Both constructs were examined by restriction enzyme digestion and verified by dideoxynucleotide sequencing.

[300] As seen in Figure 19 both the V<sub>H</sub> and the V<sub>L</sub> domains are precisely fused on their 5' ends to a gene segment encoding a leader peptide sequence (SEQ ID NO: 23 and SEQ ID NO: 24) as marked. The V<sub>H</sub> and the V<sub>L</sub> domains are ligated via *Hind* III/*Bam*HI sites into expression vector pG1D105, which contains a cDNA version of the human  $\gamma$ 1 constant region gene, and pKN100, which contains a cDNA version of the human  $\kappa$  chain constant region gene, respectively. In each case, expression is under control of the HCMVi promoter and terminated by an artificial termination sequence. The light and the heavy chain complementarily determining region (CDR) residues, defined according the hypervariable sequence definition of Kabat et al., are underlined and labeled CDR-H1 to H3 and CDR-L1 to L3, respectively. CDR-H1 (SEQ ID NO: 1 and SEQ ID NO: 9); CDR-H2 (SEQ ID NO: 2 and SEQ ID NO: 10); CDR-H3 (SEQ ID NO: 3 and SEQ ID NO: 11); CDR-L1 (SEQ ID NO: 4 and SEQ ID NO: 12); CDR-L2 (SEQ ID NO: 5 and SEQ ID NO: 13); CDR-L3 (SEQ ID NO: 6 and SEQ ID NO: 14).

[301] *Example IX-3(d). IgG expression and purification.*

[302] COS cells were co-transfected with equal amounts of c-p1C11-L and c-p1C11-H plasmids for transient IgG expression. Subconfluent COS cells grown in DMEM / 10% FCS in 150 mm culture dishes were rinsed once with 20 ml of DMEM containing 40 mM Tris (pH 7.4), followed by incubation at 37°C for 4.5 h with 4 ml of DMEM / DEAE-Dextran / DNA mixture (DMEM containing 40 mM Tris, 0.4 mg/ml of DEAE-Dextran (Sigma), and 20  $\mu$ g each of c-p1C11-L and c-p1C11 -H plasmids). The cells were incubated at 37°C for 1 h with 4 ml of DMEM / 2% FCS containing 100 nM of chloroquine (Sigma), followed by incubation with 1.5 ml of 20% glycerol / PBS at room temperature for 1 min. The cells were washed twice with DMEM / 5% FCS and incubated in 20 ml of the same medium at 37°C overnight. The cell culture medium was changed to serum-free DMEM / HEPES after the cells were washed twice with plain DMEM. The cell culture supernatant was collected at 48 h and 120 h after the transfection. The chimeric IgG was purified from the pooled supernatant by affinity chromatography using Protein G column following the protocol described by the manufacturer (Pharmacia Biotech). The IgG-containing fractions were pooled, buffer exchanged into PBS and

concentrated using Centricon 10 concentrators (Amicon Corp., Beverly, MA). The purity of the IgG was analyzed by SDS-PAGE. The concentration of purified antibody was determined by ELISA using goat anti-human  $\gamma$  chain specific antibody as the capture agent and HRP-conjugated goat anti-human  $\kappa$  chain antibody as the detection agent. Standard curve was calibrated using a clinical grade antibody, C225.

[303] After affinity purification by Protein G, a single protein band of ~150 kD was seen in SDS-PAGE. Western blot analysis using HRP-conjugated anti-human IgG1 Fc specific antibody confirmed the presence of human IgG Fc portion in the purified protein (not shown).

[304] The results of the ELISA show that c-p1C11 binds more efficiently to immobilized KDR (VEGFR-2) than the parent scFv (Fig. 20).

[305] *Example IX-4. Assays and Analysis*

[306] *Example IX-4(a). FACS analysis.*

[307] Early passage HUVEC cells were grown in growth factor-depleted EBM-2 medium overnight to induce the expression of KDR (VEGFR-2). The cells were harvested and washed three times with PBS, incubated with c-p1C11 IgG (5  $\mu$ g/ml) for 1 h at 4°C, followed by incubation with a FITC labeled rabbit anti-human Fc antibody (Capper, Organon Teknika Corp., West Chester, PA) for an additional 60 min. The cells were washed and analyzed by a flow cytometer (Model EPICS®, Coulter Corp., Edison, NJ).

[308] Figure 21 is a graph showing the FACS analysis of c-p1C11 binding to KDR (VEGFR-2)-expressing HUVEC. As previously seen with the parent scFv p1C11, c-p1C11 binds specifically to KDR (VEGFR-2) expressed on early passage HUVEC.

[309] *Example IX-4(b). Quantitative KDR (VEGFR-2) binding assay.*

[310] Various amounts of antibodies were added to KDR (VEGFR-2)-coated 96-well Maxi-sorp microtiter plates (Nunc. Denmark) and incubated at room temperature for 1 h, after which the plates were washed 3 times with PBS containing 0.1% Tween-20. The plates were then incubated at RT for 1 h with 100 u1 of mouse anti-E tag antibody-HRP

conjugate (Phannacia Biotech) for the scFv, or rabbit anti-human IgG Fc specific antibody-HRP conjugate (Cappel, Organon Teknika Corp.) for the chimeric IgG. The plates were washed 5 times, TMB peroxidase substrate (KPL, Gaithersburg, MD) added, and the OD at 450 nm read using a microplate reader (Molecular Device, Sunnyvale, CA).

[311] Figure 20 is a graph showing the direct binding of antibodies to immobilized KDR (VEGFR-2). C-p1C11 is shown to bind more efficiently to immobilized KDR (VEGFR-2) receptor than the parent scFv.

[312] *Example IX-4(c). BIA core analysis.*

[313] The binding kinetics of antibodies to KDR (VEGFR-2) were measured using BIAcore biosensor (Pharmacia Biosensor). KDR (VEGFR-2)-AP fusion protein was immobilized onto a sensor chip, and antibodies or VEGF were injected at concentrations ranging from 25 nM to 200 nM. Sensorgrams were obtained at each concentration and were evaluated using a program, BIA Evaluation 2.0, to determine the rate constants *kon* and *koff*. K<sub>d</sub> was calculated as the ratio of rate constants *koff/kon*.

[314] BIAcore analysis reveals that c-plC11 bind to KDR (VEGFR-2) with higher affinity than the parent scFv (Table 2). The K<sub>d</sub> of c-p1C11 is 0.82 nM, compared to 2.1 nM for the scFv. The increased affinity of c-p1C11 is mainly due to a slower dissociation rate (*koff*) of the bivalent chimeric IgG. It is important to note that the affinity (K<sub>d</sub>) of c-p1C11 for binding to KDR (VEGFR-2) is similar to that of the natural ligand VEGF for binding to KDR (VEGFR-2), which is 0.93 nM as determined in our BIAcore analysis (Table 2).

[315] *Example IX-4(d). Competitive VEGF binding assay.*

[316] In the first assay, various amounts of antibodies were mixed with a fixed amount of KDR (VEGFR-2)-AP (50 ng) and incubated at room temperature for 1 h. The mixtures were then transferred to 96-well microtiter plates coated with VEGF<sub>165</sub> (200 ng/well) and incubated at room temperature for an additional 2 h, after which the plates were washed 5 times and the substrate for AP (p-nitrophenyl phosphate, Sigma) was added to quantify the bound KDR (VEGFR-2)-AP molecules. EC<sub>50</sub>, i.e., the antibody concentration required for 50% inhibition of KDR (VEGFR-2) binding to VEGF, was then calculated.

[317] Figure 22 shows that c-p1C11 block KDR (VEGFR-2) receptor from binding to immobilized VEGF in a dose-dependent manner. The chimeric antibody is more potent in blocking VEGF-KDR (VEGFR-2) interaction with an  $IC_{50}$  (i.e., the antibody concentrations required to inhibit 50% of KDR (VEGFR-2) from binding to VEGF) of 0.8 nM, compared to that of 2.0 nM for the scFv. The control scFv p2A6 also binds KDR (VEGFR-2) (Fig. 20) but does not block VEGF-KDR (VEGFR-2) interaction (Fig. 22).

[318] In the second assay, various amounts of c-p1C11 antibody or cold VEGF<sub>165</sub> protein were mixed with a fixed amount of <sup>125</sup>I labeled VEGF<sub>165</sub> and added to 96-well microtiter plates coated with KDR (VEGFR-2) receptor. The plates were incubated at room temperature for 2h, washed 5 times and the amounts of radiolabeled VEGF<sub>165</sub> that bound to immobilized KDR (VEGFR-2) receptor were counted. Concentrations of c-p1C11 and cold VEGF<sub>165</sub> required to block 50% of binding of the radiolabeled VEGF to immobilized KDR (VEGFR-2) receptor were determined.

[319] The results of the inhibition of binding of radiolabeled VEGF<sub>165</sub> is shown in Figure 23. The data shown are the means of triplicate determinations. c-p1C11 is shown to efficiently compete with <sup>125</sup>I labeled VEGF for binding to immobilized KDR (VEGFR-2) receptor in a dose-dependent manner. As expected, C225, a chimeric antibody directed against EGF receptor does not bind to KDR (VEGFR-2) receptor or block VEGF-KDR (VEGFR-2) interaction (not shown).

[320] *Example IX-4(e). Phosphorylation assay.*

[321] Subconfluent HUVEC cells were grown in growth factor depleted EBM-2 medium for 24 to 48h prior to experimentation. After pretreatment with 50 nM sodium orthovanadate for 30 min, the cells were incubated in the presence or absence of antibodies for 15 min, followed by stimulation with 20 ng/ml of VEGF<sub>165</sub>, or 10 ng/ml of FGF at room temperature for an additional 15 min. The cells were then lysed in lysis buffer (50 mM Tris, 150 mM NaCl, 1% NP-40, 2 mM EDTA, 0.25% sodium deoxycholate, 1 mM PMSF, 1 µg/ml leupeptin, 1 µg/ml pepstatin, 10 µg/ml aprotinin, pH 7.5) and the cell lysate used for both the KDR (VEGFR-2) and MAP kinase phosphorylation assays. The KDR (VEGFR-2) receptor was immunoprecipitated from the cell lysates with Protein A Sepharose beads (Santa Cruz Biotechnology, Inc., CA) coupled to an anti-KDR (VEGFR-2) antibody, Mab 4.13 (ImClone Systems). Proteins were

resolved with SDS-PAGE and subjected to Western blot analysis. To detect KDR (VEGFR-2) phosphorylation, blots were probed with an antiphosphotyrosine Mab, PY20 (ICN Biomedicals, Inc. Aurora, OH). For the MAP kinase activity assay, cell lysates were resolved with SDS-PAGE followed by Western blot analysis using a phospho-specific MAP kinase antibody (New England BioLabs, Beverly, MA). All signals were detected using ECL (Amersham, Arlington Heights, IL). In both assays, the blots were reprobed with a polyclonal anti-KDR (VEGFR-2) antibody (ImClone Systems) to assure that equal amount of protein was loaded in each lane of SDS-PAGE gels.

- [322] C-p1C11 effectively inhibits VEGF-stimulated phosphorylation of KDR (VEGFR-2) receptor and activation of p44/p42 MAP kinases. In contrast, C225 does not show any inhibition of VEGF-stimulated activation of KDR (VEGFR-2) receptor and MAP kinases. Neither c-p1C11, nor C225 alone has any effects on the activity of KDR (VEGFR-2) receptor and p44/p42 MAP kinases. As previously seen with the scFv p1C11, c-p1C11 does not inhibit FGF-stimulated activation of p44/p42 MAP kinases (not shown). Furthermore, neither scFv p2A6, nor the chimeric IgG form of p2A6 (c-p2A6), inhibits VEGF-stimulated activation of KDR (VEGFR-2) receptor and MAP kinases (not shown).

- [323] *Example IX-4(f). Anti-mitogenic assay.*

- [324] The effect of anti-KDR (VEGFR-2) antibodies on VEGF-stimulated mitogenesis of human endothelial cells was determined with a [<sup>3</sup>H]-TdR DNA incorporation assay using HUVEC. HUVEC (5 x 10<sup>3</sup> cells/well) were plated into 96-well tissue culture plates in 200 µl of EBM-2 medium without VEGF, bFGF or EGF and incubated at 37°C for 72 h. Various amounts of antibodies were added to duplicate wells and pre-incubated at 37°C for 1 hour, after which VEGF<sub>165</sub> was added to a final concentration of 16 ng/ml. After 18 hours of incubation, 0.25 µCi of [<sup>3</sup>H]-TdR was added to each well and incubated for an additional 4 hours. DNA incorporated radioactivity was determined with a scintillation counter. The data shown in Figure 24 are representative of at least three separate experiments.

- [325] Both c-p1C11 and scFv p1C11 effectively inhibit mitogenesis of HUVEC stimulated by VEGF (Fig. 24). C-p1C11 is a stronger inhibitor of VEGF-induced mitogenesis of HUVEC than the parent scFv. The antibody concentrations required to inhibit 50% of VEGF-induced mitogenesis of HUVEC are 0.8 nM for c-p1C11 and 6 nM

for the scFv, respectively. As expected, scFv p2A6 does not show any inhibitory effect on VEGF-stimulated endothelial cell proliferation.

[326] The invention as claimed is enabled in accordance with the above specification and readily available references and starting materials. Nevertheless, Applicants have deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Md., 20852 USA (ATCC) the hybridoma cell lines that produce the monoclonal antibodies listed below:

[327] Hybridoma cell line DC-101 producing rat anti-mouse flk-1 (VEGFR-2) monoclonal antibody deposited on January 26, 1994 (ATCC Accession Number HB 11534).

[328] Hybridoma cell line M25.18A1 producing mouse anti-mouse flk-1 (VEGFR-2) monoclonal antibody Mab 25 deposited on July 19, 1996 (ATCC Accession Number HB 12152).

[329] Hybridoma cell line M73.24 producing mouse anti-mouse flk-1 (VEGFR-2) monoclonal antibody Mab 73 deposited on July 19, 1996 (ATCC Accession Number HB 12153).

[330] These deposits were made under the provisions of the Budapest Treaty on the International Recognition of the Deposit of Microorganisms for the Purposes of Patent Procedure and the regulations thereunder (Budapest Treaty). This assures maintenance of a viable culture for 30 years from date of deposit. The organism will be made available by ATCC under the terms of the Budapest Treaty, and subject to an agreement between Applicants and ATCC, which assures unrestricted availability upon issuance of the pertinent U.S. patent. Availability of the deposited strains is not to be construed as a license to practice the invention in contravention of the rights granted under the authority of any government in accordance with its patent laws.

TABLE 1 KDR (VEGFR-2) – BINDING ANALYSIS OF ANTI-KDR (VEGFR-2) ScFv ANTIBODIES

ScFv Clone	KDR Binding <sup>1</sup>	VEGF Binding <sup>2</sup>	Binding Kinetics <sup>3</sup>		
	(ED <sub>50</sub> nM)	(IC <sub>50</sub> nM)	<i>kon</i> (10 <sup>5</sup> M <sup>-1</sup> s <sup>-1</sup> )	<i>koff</i> (10 <sup>-4</sup> s <sup>-1</sup> )	<i>Kd</i> (10 <sup>-9</sup> M)
P1C11	Yes (0.3)	Yes (3.0)	1.1	2.3	2.1
P1F12	Yes (1.0)	Yes (15)	0.24	1.4	5.9
P2A6	Yes (5.0)	No (>300)	4.1	46.1	11.2
P2A7	No (N/A)	No (>300)	N/A	N/A	N/A

1. Determined by direct binding ELISA, numbers in the parenthesis represent the scFv concentrations that give 50% of maximum binding;

2. Determined by competitive VEGF blocking ELISA, numbers in the parenthesis represent the scFv concentrations required for 50% inhibition of KDR binding to immobilized VEGF;

3. Determined by BIAcore analysis. NA = not applicable.

TABLE 2. BINDING KINETICS OF P1C11 SCFV AND C-P1C11 TO KDR (VEGFR-2) RECEPTOR.\*

Antibody	<i>kon</i> (10 <sup>5</sup> M <sup>-1</sup> s <sup>-1</sup> )	<i>koff</i> (10 <sup>-4</sup> s <sup>-1</sup> )	<i>Kd</i> (10 <sup>-9</sup> M)
p1C11 scFv	1.11	2.27	2.1
c-p1C11	0.63	0.52	0.82
VEGF	1.87	1.81	0.93

\*All rates are determined by surface plasmon resonance using BIAcore system, and are mean of at least three separate determinations.

### [331] EXAMPLE X

[332] The present example demonstrates production of a VEGFR antagonist, namely, an anti-flt-1 (anti-VEGFR-1) monoclonal antibody, MF-1.

[333] The rat anti-VEGFR-1 monoclonal antibody was developed through a standard hybridoma technique. Eight weeks old rats were primed intraperitoneally (i.p.) with 100 µg of VEGFR-1 Fc (constant region) recombinant protein (R&D Systems, Minneapolis, MN) mixed with complete Freund's adjuvant. Then, the rats were boosted three times prior to fusion with the same protein mixed with incomplete Freund's adjuvant.

[334] Hybridoma cells were generated by fusing myeloma cells P3x63Ag8.653 with spleen cells and bone marrow cells from immunized rats. Anti-VEGFR-1 specific clones were selected using VEGFR-1 alkaline phosphatase (AP) recombinant protein in ELISA-based binding and blocking assays. Positive clones were subcloned by limiting dilution.

- [335] Anti-VEGFR-1 monoclonal antibodies (mAbs) from hybridomas were obtained via continuous feed fermentation in serum-free medium. The mAbs were purified from serum-free conditioned media by affinity chromatography using Gamma-bind protein G-Sepharose. The mAbs used in *in vivo* studies were tested for endotoxin using the Pyrogen plus® Limulus Amebocyte Lysate kit (BioWhittaker, Walkersville, MD). All antibody preparations used in animal studies contained  $\leq 1.25$  EU/ml of endotoxin. Anti-VEGFR-1 polyclonal antibodies were generated from recombinant VEGFR-1 AP protein immunized rabbit and purified by Gamma-bind protein G column (Amersham Pharmacia Biotech, Uppsala, Sweden).
- [336] The immunochemical properties of anti-VEGFR-1 mAbs were characterized in ELISA-based binding and blocking assays as well as BIAcore analysis for affinity. Binding assays were performed by coating 96-well microtiter plates (Falcon Flexible plate, Becton Dickinson, Bedford, MA) with 50 ng/well VEGFR-1 AP or VEGFR-2 AP protein overnight at 4 °C. Wells were blocked by adding 200  $\mu$ l of phosphate-buffered saline containing 5% bovine serum, 0.05% Tween 20 (blocking buffer) and incubating for 2 hrs at room temperature (RT). Wells were then washed (5x) and incubated for 1 hr at RT with various concentrations of mAbs at 50  $\mu$ l diluted in blocking buffer. Wells were again washed (5x) and incubated with 50  $\mu$ l of goat anti-rat IgG-HRP (BioSource International, Camarillo, CA) for 1 hr at RT. Wells were washed (5x) for a final time and then incubated with 50  $\mu$ l of 3,3', 5,5'-tetra-methylbenzidine (TMB) substrate (Kirkegaard and Perry Lab Inc., Gaithersburg, MD) for 15 mins at RT. The reaction was stopped by adding 50  $\mu$ l of 1 M Phosphoric Acid ( $H_3PO_4$ ) and wells read at 450 nm on a microtiter plate reader.
- [337] For VEGFR-1/VEGF or PlGF blocking assays, wells were coated with 100 ng of VEGF or PlGF (R & D Systems, Minneapolis, MN) overnight at 4°C. Wells are blocked as described above and then incubated for 1 hr at RT with 100 ng of VEGFR-1 AP that had been preincubated for 1 hr with various concentrations of mAb. Wells were washed and incubated with p-nitrophenyl phosphate (PNPP, Sigma, St. Louis, MO). Color was developed for 30 mins at RT and was then read at 405 nm on a microtiter plate reader.
- [338] The binding kinetics of anti-VEGFR-1 mAbs to VEGFR-1 was determined using BIAcore biosensor (Pharmacia Biosensor). VEGFR-1 Fc fusion protein was immobilized

onto a sensor chip and the mAbs was injected at concentrations ranging from 3.125 nM to 50 nM. Sensorgrams were obtained at each concentration and were evaluated using the program, BIA Evaluation 2.0, to determine the ratio of rate constant  $k_{on}/k_{off}$  for  $K_d$  value.

[339] *EXAMPLE XI*

[340] The present example demonstrates the inhibition of tumor growth after administration of an epidermal growth factor receptor (EGFR) antagonist, the monoclonal antibody ERBITUX<sup>TM</sup> (also known as IMC-C225 or C225), and a therapeutically effective amount of a vascular endothelial growth factor receptor (VEGFR) antagonist, the anti-VEGFR-1 monoclonal antibody DC101.

[341] Antibodies for immunohistochemical analysis and antiangiogenic therapy were obtained as follows: rat anti-mouse CD31/PECAM-1 antibody from Pharmingen (San Diego, CA); mouse anti-proliferating cell nuclear antigen (PCNA) clone PC10 DAKO A/S from DAKO Corp. (Carpinteria, CA); peroxidase-conjugated goat anti-rat immunoglobulin (IgG) (H+L) and Texas Red-conjugated goat anti-rat IgG from Jackson Research Laboratories (West Grove, PA); peroxidase-conjugated rat antimouse IgG2a from Serotec Harlan Bioproducts for Science, Inc. (Indianapolis, IN); and rat anti-mouse VEGF receptor-2 monoclonal antibody (Prewett et al., 1999; Witte et al., 1998) and chimeric anti-human EGF receptor monoclonal antibody (Goldstein et al., 1995) from ImClone Systems, Inc. (New York, NY) (Prewett et al., 1999; Witte et al., 1998).

[342] Eight-week-old male athymic nude mice (National Cancer Institute, Animal Production Area, Frederick, MD), were given an intraperitoneal injection of  $1.0 \times 10^6$  KM12L4 colon cancer cells in 500  $\mu$ l of HBSS with a 30-gauge needle attached to a 1-ml syringe. Mice were then randomized into one of four treatment groups (10 mice per group): control, DC101, C225, or DC101 and C225. All animal studies were conducted under guidelines approved by the Animal Care and Use Committee of MD Anderson Cancer Center and UKCCCR, 1998).

[343] Beginning 3 days after injection of tumor cells, mice were given intraperitoneal (i.p.) injections every third day of either control vehicle (phosphate buffered saline [PBS]), DC101 (0.8 mg), C225 (1.0 mg), or DC101 (0.8 mg) and C225 (1.0 mg) (each injection

given in a 700- $\mu$ l total volume) with a 27-gauge needle attached to a 1-ml syringe. Mice were weighed weekly. Mice were killed when the control group became moribund (about 30 days after therapy began), and complete necropsies were performed, tumor burden was quantified, and tumors were harvested and analysed as described below.

- [344] For each mouse, necropsy was performed, the size of peritoneal tumors was measured with calipers, the mean tumor size was determined, and representative lesions were excised. The extent of ascites was assessed by an investigator who was unaware of the treatment-group assignment as follows: grade 0, no ascites; grade 1, small ascites; grade 2, moderate ascites; grade 3, large ascites; grade 4, massive ascites with tense abdomen (Aparicio et al., 1999). Tumor sections were either embedded in OCT compound and frozen at -70°C or fixed in formalin and then embedded in paraffin.
- [345] Tissue sections were treated by standard deparaffinization (for formalin-fixed and paraffin-embedded tissues) or by fixation in acetone and chloroform (for tissues frozen in OCT), and immuno-histochemical analyses were performed as described previously (Shaheen et al., 1999). Briefly, endogenous peroxidases were blocked with 3% H<sub>2</sub>O<sub>2</sub> in methanol, and the slides were washed with PBS, incubated for 20 min in protein-blocking solution (PBS supplemented with 1% normal goat serum and 5% normal horse serum), and incubated overnight at 4°C with primary antibodies against CD31 or PCNA. Then, the slides were washed, incubated with protein-blocking solution, incubated for 1 h at room temperature with peroxidase-conjugated secondary antibodies, washed incubated with DAB, washed, counterstained with haematoxylin, washed, and mounted with Universal Mount and dried on a hot plate at 56°C. Frozen sections to be stained for CD31 were incubated with a secondary antibody conjugated to Texas Red (red fluorescence) instead of the peroxidase-conjugated antibody. Omission of the primary antibody served as negative control.
- [346] Terminal deoxynucleotidyl transferase-(TdT)-mediated dUTP nick-end labelling (TUNEL) staining was performed according to the manufacturer's protocol. Briefly, the sections were fixed with 4% methanol-free paraformaldehyde, washed, permeabilized with 0.2% Triton X-100, washed, incubated with the kit's equilibration buffer, incubated with a reaction mix containing equilibration buffer, nucleotide mix, and the TdT enzyme at 37°C for 1 h, incubated for 15 min at room temperature with 2x standard saline citrate to stop

the TdT reaction, washed, stained with DAPI mount (to visualize the nuclei), and glass coverslips were applied.

[347] Numbers of tumor vessels and PCNA-positive cells were evaluated by light microscopy (counted in five random  $0.159\text{-mm}^2$  fields at 100x magnification), imaged digitally, and processed with Optimas Image Analysis software (Biscan, Edmond, WA). Apoptotic cells were visualized with immunofluorescence as follows. Sections were digitally imaged and processed with Adobe Photoshop software (Adobe Systems, Mountain View, CA). CD31-positive endothelial cells (ECs) were detected by localized red fluorescence by using a rhodamine filter. Apoptosis was determined by localized green fluorescence for tumor cells (TCs) or green with red fluorescence for ECs by using a fluorescence filter. Nuclei were detected by the blue fluorescence of the DAPI with its filter. Cells were counted in five consecutive, non-overlapping fields  $0.011\text{-mm}^2$  fields per slide at 400x magnification with the first field selected at random in a non-necrotic portion of the tumor. The percentages of apoptotic ECs and TCs per field were then calculated as [% apoptotic cells = (number of apoptotic cells/total number of cells)x100].

[348] Measuring the diameter for the peritoneal lesions 30 days after therapy began was used to assess gross tumor burden. The mean peritoneal tumor size was smaller in the DC101 group (50.3 % smaller) and in the combination DC101+C225 group (66.7 % smaller) than in the control group. Although 100% of the control and C225 mice had peritoneal disease at the end of the study, only 10 % of the DC101 mice and 30 % of the combination therapy mice showed no evidence of disease. Finally, the mean ascites grade was lower for both the DC101 (66.7 % lower) and combination therapy groups (100 % lower) than for the control mice. Moreover, the DC101+C225 group had significantly less ascites than did the DC101 group; virtually no ascites was found in the combination group.

[349] Tumor cells were stained for PCNA by immunohistochemical analysis to assess tumor cell proliferation. Peritoneal tumors were smaller in the DC101 (50.3 % smaller) and DC101+C225 groups (66.7 % smaller) than in the control group.

[350] Section of the peritoneal lesions for mice were also stained for CD31 immunofluorescence to detect the number of ECs as a measure of angiogenesis. Significantly fewer ECs was observed in the DC101 and DC101+C225 groups than in the control group.

- [351] Immunofluorescent TUNEL staining, with and without concurrent staining for CD31, was performed on sections of peritoneal metastases to quantify TC and EC apoptosis. More apoptotic TCs and ECs were observed in the DC101 group (14.3-fold more for TCs and 226-fold more for ECs) and DC101+C225 group (23.6-fold more for TCs and 331-fold more for ECs) than in the control group, and more apoptotic TCs and ECs were observed in the DC101+C225 group than in the DC101 group alone ((1.7-fold more for TCs and 1.46-fold more for ECs).
- [352] *EXAMPLE XII. PRODUCTION OF HUMAN FAB*
- [353] *Example XII(a). Proteins and Cell Lines.*
- [354] Primary-cultured HUVEC were obtained from Dr. S. Rafii at Cornell Medical Center, New York, and maintained in EBM-2 medium (Clonetics, Walkersville, MD) at 37°C, 5% CO<sub>2</sub>. The soluble fusion proteins, KDR (VEGFR-2)-AP, its immunoglobulin (Ig) domain-deletion variants, and Flk-1-AP, were expressed in stably transfected NIH 3T3 and purified from cell culture supernatants by affinity chromatography using immobilized monoclonal antibody to AP as described by Lu et al., *J. Biol. Chem.* 275: 14321-30 (2000). VEGF<sub>165</sub> protein was expressed in baculovirus and purified following the procedures described in Zhu et al., *Cancer Res.* 58: 3209-14 (1998). The leukemia cell lines, HL60 and HEL, were maintained in RPMI containing 10% fetal calf serum.
- [355] *Example XII(b). Phage ELISA*
- [356] Individual TG1 clones were picked and grown at 37°C in 96 well plates and rescued with M13K07 helper phage as described above. The amplified phage preparation was blocked with 1/6 volume of 18% milk/PBS at RT for 1 h and added to Maxi-sorp 96-well microtiter plates (Nunc) coated with KDR (VEGFR-2)-AP or AP (1 µg/ml x 100 µl). After incubation at RT for 1 h the plates were washed 3 times with PBST and incubated with a rabbit anti-M13 phage-HRP conjugate (Amersham Pharmacia Biotech, Piscataway, NJ). The plates were washed 5 times, TMB peroxidase substrate (KPL, Gaithersburg, MD) added, and the absorbance at 450 nm read using a microplate reader (Molecular Devices, Sunnyvale, CA).

[357] *Example XII(c). DNA BstNI pattern analysis and nucleotide sequencing.*

[358] The diversity of the anti-KDR (VEGFR-2) Fab clones after each round of selection was analyzed by restriction enzyme digestion patterns (i.e., DNA fingerprints). The Fab gene insert of individual clones was PCR amplified using primers: PUC19 reverse, 5' AGCGGATAACAATTTACACAGG 3'; and fdtet seq, 5' GTCGTCTTTCCAGACGTTAGT 3'. The amplified product was digested with a frequent-cutting enzyme, *BstNI*, and analyzed on a 3% agarose gel. DNA sequences of representative clones from each digestion pattern were determined by dideoxynucleotide sequencing.

[359] *Example XII(d). Expression and purification of soluble Fab fragments.*

[360] Plasmids of individual clones were used to transform a nonsuppressor *E. coli* host HB2151. Expression of the Fab fragments in HB2151 was induced by culturing the cells in 2YTA medium containing 1 mM isopropyl-1-thio- $\beta$ -D-galactopyranoside (IPTG, Sigma) at 30°C. A periplasmic extract of the cells was prepared by resuspending the cell pellet in 25 mM Tris (pH 7.5) containing 20% (w/v) sucrose, 200 mM NaCl, 1 mM EDTA and 0.1 mM PMSF, followed by incubation at 4°C with gentle shaking for 1 h. After centrifugation at 15,000 rpm for 15 min, the soluble Fab protein was purified from the supernatant by affinity chromatography using a Protein G column followed the manufacturer's protocol (Amersham Pharmacia Biotech).

[361] *Example XII(e). Selection of human anti-KDR (VEGFR-2) Fab from phage display library.*

[362] A large human Fab phage display library containing  $3.7 \times 10^{10}$  clones (DeHaard et al., *J. Biol. Chem.* 274: 18218-30 (1999)) was used for the selection. The library consists of PCR-amplified antibody variable light chain genes and variable heavy chain genes fused to human constant light chain genes ( $\kappa$  and  $\lambda$ ) and DNA encoding the IgG1 heavy chain C<sub>H</sub>1 domain, respectively. Both heavy and light chain constructs are preceded by a signal sequence - *pelB* for the light chain and gene III signal sequence for the heavy chain. Heavy chain constructs further encode a portion of the gene III protein for phage display, a hexahistidine tag, and an 11 amino-acid-long c-myc tag, followed by an amber codon (TAG). The hexahistidine and c-myc tags can be used for purification or detection. The amber codon allows for phage display using suppressor hosts (such as TG1 cells) or

production of Fab fragments in soluble form when transformed into a nonsuppressor host (such as HB2151 cells).

- [363] The library stock was grown to log phase, rescued with M13-KO7 helper phage and amplified overnight in 2YTAK medium (2YT containing 100  $\mu\text{g/ml}$  of ampicillin and 50  $\mu\text{g/ml}$  of kanamycin) at 30°C. The phage preparation was precipitated in 4% PEG/0.5M NaCl, resuspended in 3% fat-free milk/PBS containing 500  $\mu\text{g/ml}$  of AP protein and incubated at 37°C for 1 h to capture phage displaying anti-AP Fab fragments and to block other nonspecific binding.
- [364] KDR (VEGFR-2)-AP (10  $\mu\text{g/ml}$  in PBS) coated Maxisorp Star tubes (Nunc, Roskilde, Denmark) were first blocked with 3% milk/PBS at 37°C for 1 h, and then incubated with the phage preparation at RT for 1 h. The tubes were washed 10 times with PBST (PBS containing 0.1% Tween-20) followed by 10 times with PBS. Bound phage were eluted at RT for 10 min with 1 ml of a freshly prepared solution of 100 mM triethylamine (Sigma, St. Louis, MO). The eluted phage were incubated with 10 ml of mid-log phase TG1 cells at 37°C for 30 min stationary and 30 min shaking. The infected TG1 cells were pelleted and plated onto several large 2YTAG plates and incubated overnight at 30°C. All the colonies grown on the plates were scraped into 3 to 5 ml of 2YTA medium, mixed with glycerol (10% final concentration), aliquoted and stored at -70°C. For the next round selection, 100  $\mu\text{l}$  of the phage stock was added to 25 ml of 2YTAG medium and grown to mid-log phase. The culture was rescued with M13K07 helper phage, amplified, precipitated, and used for selection followed the procedure described above, with reduced concentrations of KDR (VEGFR-2)-AP immobilized on the immunotube and increased number of washes after the binding process.
- [365] A total of three rounds of selection were performed on immobilized KDR (VEGFR-2), with varying protein concentrations and number of washings after the initial binding process. After each round selection, 93 clones were randomly picked and tested by phage ELISA for binding to KDR (VEGFR-2). Seventy out of the 93 clones (75%) picked after the second selection, and greater than 90% of the recovered clones after the third selection were positive in KDR (VEGFR-2) binding, suggesting a high efficiency of the selection process. DNA segments encoding the Fab from all the 70 binders identified in the second selection were amplified, digested with *Bst*NI, and compared for fingerprint

patterns. A total of 42 different patterns were observed, indicating an excellent diversity of the isolated anti-KDR (VEGFR-2) Fab. Cross-reactivity examination demonstrated that 19 out of the 42 antibodies were specific FLT-1 (VEGFR-1)-binders, whereas the rest 23 antibodies bound to both KDR (VEGFR-2) and its murine homologue, Flk-1. Further selection was achieved with a competitive VEGF-binding assay in which the binding of soluble KDR (VEGFR-2) to immobilized VEGF in the presence or absence of the anti-KDR (VEGFR-2) Fab fragments was determined. The assay identified four Fab clones that were capable of blocking the binding between VEGF and KDR (VEGFR-2). Three were KDR (VEGFR-2)-specific binders and one cross-reacted with Flk-1. DNA fingerprinting and sequencing analysis confirmed that all four KDR (VEGFR-2)/VEGF blocking antibodies were different with unique DNA and amino acid sequences.

[366] The amino acid sequences for CDR1, CDR2 and CDR3 of V<sub>H</sub> and V<sub>L</sub> for the four clones are given in Table 3.

**TABLE 3 - CDR SEQUENCES OF SELECTED KDR (VEGFR-2)-BINDING HUMAN FABS**

Clone	CDR1	CDR2	CDR3
<b>Light Chain</b>			
D2C6	RASQSVSSYLA (SEQ ID NO:1)	DSSNRAT (SEQ ID NO:2)	LQHNTFPPT (SEQ ID NO:3)
D2H2	RASQGISSRLA (SEQ ID NO:4)	AASSLQT (SEQ ID NO:5)	QQANRFPPT (SEQ ID NO:6)
D1H4	AGTTTDLTYDDLVS (SEQ ID NO:7)	DGNKRPS (SEQ ID NO:8)	NSYVSSRFYV (SEQ ID NO:9)
D1F7	SGSTSNIGTNTAN (SEQ ID NO:10)	NNNQRP (SEQ ID NO:11)	AAWDDSLNGHWV (SEQ ID NO:12)
<b>Heavy Chain</b>			
D2C6	GFTFSSYSMN (SEQ ID NO:13)	SISSSSYIYYADSVKG (SEQ ID NO:14)	VTDAFDI (SEQ ID NO:15)
D2H2	GFTFSSYSMN	SISSSSYIYYADSVKG	VTDAFDI
D1H4	GFTFSSYSMN	SISSSSYIYYADSVKG	VTDAFDI
D1F7	GGTFSSYAIS (SEQ ID NO:16)	GGIPIFGTANYAQKFQG (SEQ ID NO:17)	GYDYYDSSGVASPFDY (SEQ ID NO:18)

[367] Complete sequences for the V<sub>H</sub> and V<sub>L</sub> chains are presented in the Sequence Listing. For D1F7, the nucleotide and amino acid sequences for V<sub>H</sub> are represented by

SEQ ID NOS: 19 and 20 respectively, and the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 21 and 22.

[368] For D2C6, the nucleotide and amino acid sequences for V<sub>H</sub> are represented by SEQ ID NOS: 23 and 24 respectively, and the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 25 and 26.

[369] For D2H2, the nucleotide and amino acid sequences for V<sub>H</sub> are represented by SEQ ID NOS: 30 and 31 respectively, and the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 32 and 33.

[370] For D1H4, the nucleotide and amino acid sequences for V<sub>H</sub> are represented by SEQ ID NOS: 27 and 24 respectively, and the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 28 and 29.

[371] A second library was created combining the single heavy chain of D2C6 with a diverse population of light chains derived from the original library. Ten additional Fabs were identified, designated SA1, SA3, SB10, SB5, SC7, SD2, SD5, SF2, SF7, and 1121. The nucleotide and amino acid sequences for V<sub>L</sub> of the ten Fabs are represented as follows. For SA1, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 34 and 35. For SA3, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 36 and 37. For SB10, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 38 and 39. For SB5, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 40 and 41. For SC7, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 42 and 43. For SD2, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 44 and 45. For SD5, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 46 and 47. For SF2, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 48 and 49. For SF7, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 50 and 51. For 1121, the nucleotide and amino acid sequences for V<sub>L</sub> are represented by SEQ ID NOS: 52 and 53.

[372] The V<sub>L</sub> CDR sequences are presented in Table 4.

**TABLE 4 - LIGHT CHAIN CDR SEQUENCES OF KDR (VEGFR-2)-BINDING HUMAN FABS**

Clone	CDR1	CDR2	CDR3
SA1	TGSHSNFGAGTDV (SEQ ID NO:54)	GDSNRPS (SEQ ID NO:55)	QSYDYGLRGWV (SEQ ID NO:56)
SA3	RASQNINNYLN (SEQ ID NO:57)	AASTLQS (SEQ ID NO:58)	QQYSRYPPT (SEQ ID NO:59)
SB10	TGSSTDVGNVNYIS (SEQ ID NO:60)	DVTSRPS (SEQ ID NO:61)	NSYSATDTLV (SEQ ID NO:62)
SB5	TGQSSNIGADYDVH (SEQ ID NO:63)	GHNNRPS (SEQ ID NO:64)	QSYDSSLGLV (SEQ ID NO:65)
SC7	RASQDISWLA (SEQ ID NO:66)	AASLLQS (SEQ ID NO:67)	QQADSFPT (SEQ ID NO:68)
SD2	RASQSIKRWLA (SEQ ID NO:69)	AASTLQS (SEQ ID NO:70)	QQANSFPPT (SEQ ID NO:71)
SD5	SGSRNIGAHEVQ (SEQ ID NO:72)	GDTNRPS (SEQ ID NO:73)	QSYDTSRGPV (SEQ ID NO:74)
SF2	TGSSNIGTGVDVH (SEQ ID NO:75)	AYTNRPS (SEQ ID NO:76)	QSFDDSLNGLV (SEQ ID NO:77)
SF7	TGSHSNFGAGTDVH (SEQ ID NO:78)	GDTHRPS (SEQ ID NO:79)	QSYDYGLRGWV (SEQ ID NO:80)
1121	RASQGIDNWL (SEQ ID NO:81)	DASNLD (SEQ ID NO:82)	QQAKAFPPT (SEQ ID NO:83)

**[373] EXAMPLE XIII. ASSAYS****[374] Example XIII(a). Quantitative KDR (VEGFR-2) binding and blocking of KDR (VEGFR-2)/VEGF interaction.**

**[375]** In a direct binding assay, various amounts of soluble Fab proteins were added to KDR (VEGFR-2)-coated 96-well Maxi-sorp microtiter plates and incubated at RT for 1 h, after which the plates were washed 3 times with PBST. The plates were then incubated at RT for 1 h with 100  $\mu$ l of a rabbit anti-human Fab antibody-HRP conjugate (Jackson ImmunoResearch Laboratory Inc., West Grove, PA). The plates were washed and developed following the procedure described above for the phage ELISA. In a competitive KDR (VEGFR-2)/VEGF blocking assay, various amounts of Fab proteins were mixed with a fixed amount of KDR (VEGFR-2)-AP (100 ng) and incubated at RT for 1 h. The mixtures were then transferred to 96-well microtiter plates precoated with VEGF<sub>165</sub> (200 ng/well) and incubated at RT for an additional 2 h, after which the plates were washed 5 times and the substrate for AP (p-nitrophenyl phosphate, Sigma) was added. Absorbance at 405nm was measured to quantify the bound KDR (VEGFR-2)-AP

molecules (8).  $IC_{50}$ , *i.e.*, the Fab protein concentration required for 50% inhibition of KDR (VEGFR-2) binding to VEGF, was then calculated.

- [376] The four VEGF-blocking clones (D2C6, D2H2, D1H4, D1F7) were expressed as soluble Fab and purified from periplasmic extracts of *E. coli* by Protein G affinity chromatography. The yield of purified Fab proteins of these clones ranged from 60 to 400  $\mu\text{g}$  / liter culture. SDS-PAGE analysis of each purified Fab preparation yielded a single protein band with expected molecular size.
- [377] Clone D2C6 and D2H2 are more efficient binders, followed by clone D1H4 and D1F7. All four Fabs also block KDR (VEGFR-2) binding to immobilized VEGF. The antibody concentrations required for 50% of inhibition of KDR (VEGFR-2) binding to VEGF are approximately 2 nM for clones D2C6, D2H2, and D1H4 and 20 nM for clone D1F7. Only clone D1F7 blocks VEGF from binding to Flk-1, with an  $IC_{50}$  of approximately 15 nM.
- [378] *Example XIII (b). BIAcore analysis of the soluble scFv*
- [379] The binding kinetics of soluble Fab proteins to KDR (VEGFR-2) were measured by surface plasmon resonance using a BIAcore biosensor (Pharmacia Biosensor). KDR (VEGFR-2)-AP fusion protein was immobilized onto a sensor chip and soluble Fab proteins were injected at concentrations ranging from 1.5 nM to 100 nM. Sensorgrams were obtained at each concentration and were evaluated using a program, BIA Evaluation 2.0, to determine the rate constants  $k_{on}$  and  $k_{off}$ .  $K_d$  was calculated from the ratio of rate constants  $k_{off}/k_{on}$ .
- [380] All three KDR (VEGFR-2)-specific Fab fragments bind to immobilized receptor with  $K_d$  of 2 to 4 nM (Table 5). The cross-reactive clone, D1F7, has a  $K_d$  of 45 nM, which is about 10- to 15-fold weaker than those of the KDR (VEGFR-2)-specific clones. It is noteworthy that, although the overall  $K_d$  for the three KDR (VEGFR-2)-specific Fab fragments are similar, the individual binding kinetics, *i.e.*, the  $k_{on}$  and  $k_{off}$ , for these antibodies are quite different, *e.g.*, D2C6 possesses the fastest on-rate, while D1H4 has the slowest off-rate (Table 5).

**TABLE 5 - BINDING KINETICS OF THE FOUR  
NEUTRALIZING HUMAN ANTI-KDR (VEGFR-2) FAB  
FRAGMENTS**

Clone	$k_{on}$ ( $10^4 \text{ M}^{-1} \text{ S}^{-1}$ )	$k_{off}$ ( $10^{-4} \text{ S}^{-1}$ )	$K_d$ (nM)
Hu-2C6 Fab	$27.3 \pm 8.6^*$	$5.38 \pm 0.54$	1.97
Hu-2H2 Fab	$12.4 \pm 2.9$	$4.87 \pm 0.18$	3.93
Hu-1H4 Fab	$5.55 \pm 0.59$	$1.53 \pm 0.22$	2.76
Hu-1F7 Fab	$4.14 \pm 1.21$	$18.7 \pm 2.12$	45.2

[381] All numbers are determined by BIAcore analysis and represent the mean  $\pm$  SE from at least three separate determinations.

[382] *Example XIII(c). Binding epitope mapping*

[383] The production of KDR (VEGFR-2) extracellular Ig-like domain deletion variants has been previously described (Lu et al. (2000)). In an epitope-mapping assay, full length KDR (VEGFR-2)-AP, Ap fusions of two KDR (VEGFR-2) Ig-domain deletion variants, and Flk-1-AP were first immobilized onto a 96-well plate (Nunc) using a rabbit anti-AP antibody (DAKO-immunoglobulins, Glostrup, Denmark) as the capture reagent. The plate was then incubated with various anti-KDR (VEGFR-2) Fab proteins at RT for 1 h, followed by incubation with a rabbit anti-human Fab antibody-HRP conjugate. The plate was washed and developed as described above.

[384] The binding epitopes of the anti-KDR (VEGFR-2) Fab fragments were mapped using the full-length KDR (VEGFR-2) and two KDR (VEGFR-2) Ig domain-deletion variants. KDR (VEGFR-2)(1-3) is a KDR (VEGFR-2) variant containing the first three N-terminal Ig domains. KDR (VEGFR-2)(3) is a variant containing only the third Ig domain. Clones D2C6 and D1H4 bind equally well to KDR (VEGFR-2), KDR (VEGFR-2)(1-3) and KDR (VEGFR-2)(3), thus locating their binding epitope(s) within Ig domain 3. Clones D2H2 and D1F7 bind much more efficiently to full-length KDR (VEGFR-2) and KDR (VEGFR-2)(1-3), indicating a broader binding epitope(s) within KDR (VEGFR-2) Ig domains 1 to 3. Only clone D1F7 cross-reacts with Flk-1.

[385] *Example XIII(d). Anti-mitogenic assay.*

[386] HUVEC ( $5 \times 10^3$  cells/well) were plated onto 96-well tissue culture plates (Wallach, Inc., Gaithersburg, MD) in 200  $\mu$ l of EBM-2 medium without VEGF, basic fibroblast growth factor (bFGF) or epidermal growth factor (EGF) and incubated at 37°C for 72 h. Various amounts of Fab proteins were added to duplicate wells and pre-incubated at 37°C for 1 h, after which VEGF<sub>165</sub> was added to a final concentration of 16 ng/ml. After 18 h of incubation, 0.25  $\mu$ Ci of [<sup>3</sup>H]TdR (Amersham) was added to each well and incubated for an additional 4 h. The cells were washed once with PBS, trypsinized and harvested onto a glass filter (Printed Filtermat A, Walach) with a cell harvester (Harvester 96, MACH III, TOMTEC, Orange, CT). The membrane was washed three times with H<sub>2</sub>O and air-dried. Scintillation fluid was added and DNA incorporated radioactivity was determined on a scintillation counter (Wallach, Model 1450 Microbeta Scintillation Counter).

[387] The ability of human anti-KDR (VEGFR-2) Fab to block VEGF-stimulated mitogenic activity on HUVEC. All four human Fab fragments inhibited VEGF induced DNA synthesis in HUVEC in a dose-dependent manner. The Fab concentration that inhibited 50% (EC<sub>50</sub>) of VEGF-stimulated [<sup>3</sup>H]-TdR incorporation in HUVEC, is approximately 0.5 nM for clones D2C6 and D1H4, 0.8 nM for clone D2H2, and 15 nM for clone D1F7. Controls included VEGF only (1500 cpm) and plain medium (60 cpm). Duplicate wells were assayed. The data shown are representative of at least three separate experiments.

[388] *Example XIII(e). Leukemia migration assay.*

[389] HL60 and HEL cells were washed three times with serum-free plain RPMI 1640 medium and suspended in the medium at  $1 \times 10^6$ /ml. Aliquots of 100  $\mu$ l cell suspension were added to either 3- $\mu$ m-pore transwell inserts for HL60 cells, or 8- $\mu$ m-pore transwell inserts for HEL cells (Costar®, Corning Incorporated, Corning, NY) and incubated with the anti-KDR (VEGFR-2) Fab proteins (5  $\mu$ g/ml) for 30 min at 37°C. The inserts were then placed into the wells of 24-well plates containing 0.5 ml of serum-free RPMI 1640 with or without VEGF<sub>165</sub>. The migration was carried out at 37°C, 5% CO<sub>2</sub> for 16-18 h for HL60 cells, or for 4 h for HEL cells. Migrated cells were collected from the lower

compartments and counted with a Coulter counter (Model Z1, Coulter Electronics Ltd., Luton, England).

[390] VEGF induced migration of HL60 and HEL cells in a dose-dependent manner with maximum stimulation achieved at 200 ng/. All the anti-KDR (VEGFR-2) Fab fragments significantly inhibited VEGF-stimulated migration of HL60 and HEL cells. As a control, a Fab fragment of C225, an antibody directed against EGF receptor, did not show significant inhibitory effect in this assay.

[391] *EXAMPLE XIV. PRODUCTION OF IgG*

[392] *Example XIV(a). Construction of vectors for expression of IgG.*

[393] Separate vectors for expression of IgG light chain and heavy chains were constructed. Cloned V<sub>L</sub> genes were digested and ligated into the vector pKN100. Cloned V<sub>H</sub> genes were digested and ligated into the vector pGID105 containing the human IgG I (γ) heavy chain constant domain. Constructs were examined by restriction enzyme digestion and verified by dideoxynucleotide sequencing. In both cases, expression is under control of the HCMV promoter and terminated by an artificial termination sequence.

[394] The assembled heavy and light chain genes were then cloned into Lonza GS expression vectors pEE6.1 and pEE12.1. Heavy and light chain vectors were recombined into a single vector for stable transfection of CHO cells and NS0 cells. Transfected cells are cultured in glutamine minus medium and express antibodies at levels as high as 1g/L.

What is claimed is:

1. A method of inhibiting tumor growth comprising administering to a human a therapeutically effective amount of a vascular endothelial growth factor (VEGF) receptor antagonist and a therapeutically effective amount of an epidermal growth factor receptor (EGFR) antagonist.
2. A method of inhibiting tumor growth comprising administering to a human a therapeutically effective amount of a vascular endothelial growth factor (VEGF) receptor antagonist and radiation.
3. A method of inhibiting tumor growth comprising administering to a human a therapeutically effective amount of a vascular endothelial growth factor (VEGF) receptor antagonist and a chemotherapeutic agent.
4. The method of any of claims 1-3, wherein the tumor overexpresses VEGFR.
5. The method of any of claims 1-4, wherein the tumor is a tumor of the breast, heart, lung, small intestine, colon, spleen, kidney, bladder, head and neck, ovary, prostate, brain, pancreas, skin, bone, bone marrow, blood, thymus, uterus, testicles, cervix or liver.
6. The method of any of claims 1-5, wherein the tumor is a tumor of the colon.
7. The method of any of claims 1-5, wherein the tumor is a non-small cell lung carcinoma (NSCLC).
8. The method of any of claims 1-7, wherein the VEGF receptor antagonist is administered intravenously.
9. The method of any of claims 1-8, wherein the VEGF receptor antagonist is administered orally.
10. The method of any of claims 1-9, wherein the VEGF receptor antagonist inhibits binding of VEGFR to its ligand.
11. The method of any of claims 1-10, wherein the VEGF receptor antagonist binds VEGFR.

12. The method of any of claims 1-11, wherein the VEGF receptor antagonist comprises an antibody, or functional equivalent thereof, specific for VEGFR.

13. The method of any of claims 1-11, wherein the VEGF receptor antagonist comprises a small molecule specific for VEGFR.

14. The method of any of claims 1-13, wherein the VEGF receptor antagonist is an antagonist of *fms*-like tyrosine kinase receptor (flt-1) VEGFR-1.

15. The method of any of claims 1-14, wherein the method further comprises administering an antineoplastic agent.

16. The method of claim 15, wherein the antineoplastic agent is a chemotherapeutic agent or radiation.

17. The method of claim 15, wherein the method further comprises administering an adjuvant.

18. The method of any of claims 1-17, wherein the tumor overexpresses EGFR.

19. The method of any of claims 1, 4-18, wherein the EGFR antagonist is administered intravenously.

20. The method of any of claims 1, 4-19, wherein the EGFR antagonist is administered orally.

21. The method of any of claims 1, 4-20, wherein the EGFR antagonist inhibits binding of EGFR to its ligand.

22. The method of any of claims 1, 4-21, wherein the EGFR antagonist binds EGFR.

23. The method of any of claims 1, 4-24, wherein the EGFR antagonist inhibits binding of EGFR to ATP.

24. The method of any of claims 1, 4-22, wherein the EGFR antagonist comprises an antibody, or functional equivalent thereof, specific for EGFR.

25. The method of any of claims 1, 4-23, wherein the EGFR antagonist comprises a small molecule specific for EGFR.

26. The method of any of claims 1, 4-25, wherein the antibody comprises a constant region of a human antibody.

27. The method of claim 26, wherein the antibody is a chimeric antibody comprising a variable region of a mouse antibody.

28. The method of claim 26, wherein the antibody is a humanized antibody comprising a variable region having complementarity-determining regions (CDRs) of a mouse antibody and framework regions of a human antibody.

29. The method of claim 26, wherein the antibody is a human antibody comprising a variable region of a human antibody.

30. A method of any of claims 3-29, wherein the chemotherapeutic agent is not conjugated to the VEGF receptor antagonist.

31. A method of any of claims 3-30, wherein the chemotherapeutic agent is selected from the group consisting of cisplatin, dicarbazine, dactinomycin, mechlorethamine, streptozocin, cyclophosphamide, carmustine, lomustine, doxorubicin, daunorubicin, procarbazine, mitomycin, cytarabine, etoposide, methotrexate, 5-fluorouracil, vinblastine, vincristine, bleomycin, paclitaxel, docetaxel, aldesleukin, asparaginase, busulfan, carboplatin, cladribine, dacarbazine, floxuridine, fludarabine, hydroxyurea, ifosfamide, interferon alpha, leuprolide, megestrol, melphalan, mercaptopurine, plicamycin, mitotane, pegaspargase, pentostatin, pipobroman, plicamycin, streptozocin, tamoxifen, teniposide, testolactone, thioguanine, thiotepa, uracil mustard, vinorelbine, chlorambucil, and taxol and combinations thereof.

32. A method of any of claims 3-31, wherein the chemotherapeutic agent is selected from the group consisting of cisplatin, doxorubicin, taxol and combinations thereof.

33. A kit for inhibiting tumor growth comprising a therapeutically effective amount of an epidermal growth factor receptor (EGFR) antagonist and a therapeutically effective amount of a vascular endothelial growth factor (VEGF) receptor antagonist.

34. The kit of claim 33, wherein the EGFR antagonist comprises an antibody, or functional equivalent thereof, specific for EGFR.

35. The kit of claim 33, wherein the EGFR antagonist comprises a small molecule specific for EGFR.

36. The kit of any of claims 33-35, wherein the VEGFR antagonist comprises an antibody, or functional equivalent thereof, specific for VEGFR.

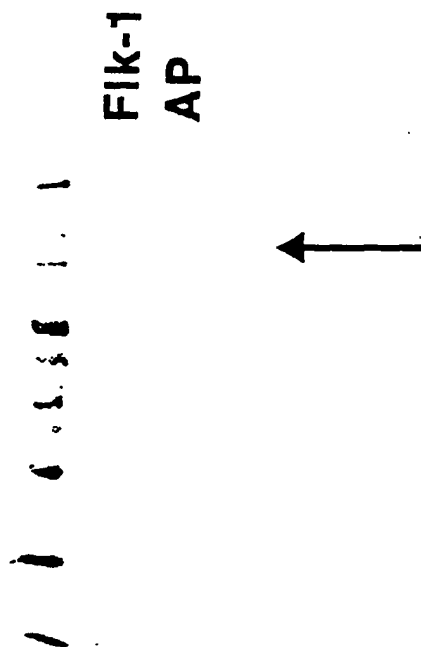
37. The kit of any of claims 33-35, wherein the VEGFR antagonist comprises a small molecule specific for VEGFR.

38. The kit of any of claims 33-37, wherein the kit further comprises an antineoplastic agent.

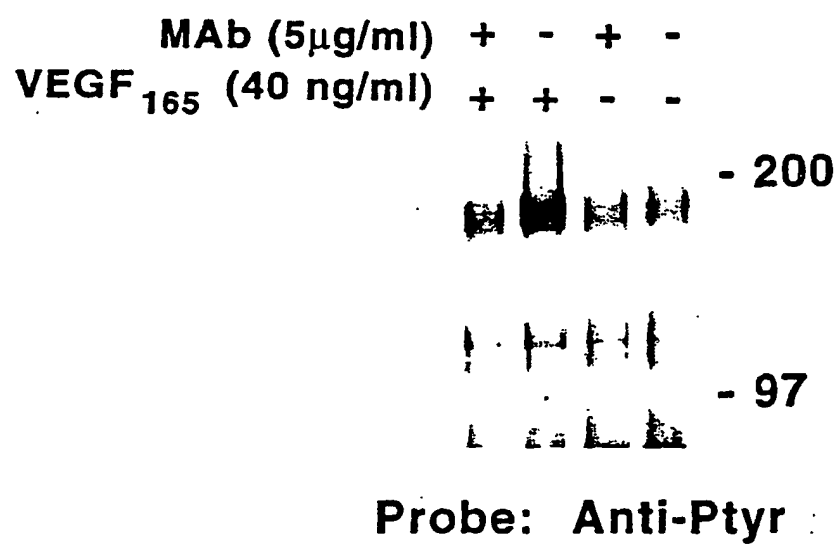
The kit of any of claims 33-38, wherein the kit further comprises an adjuvant.

# FIGURE 1

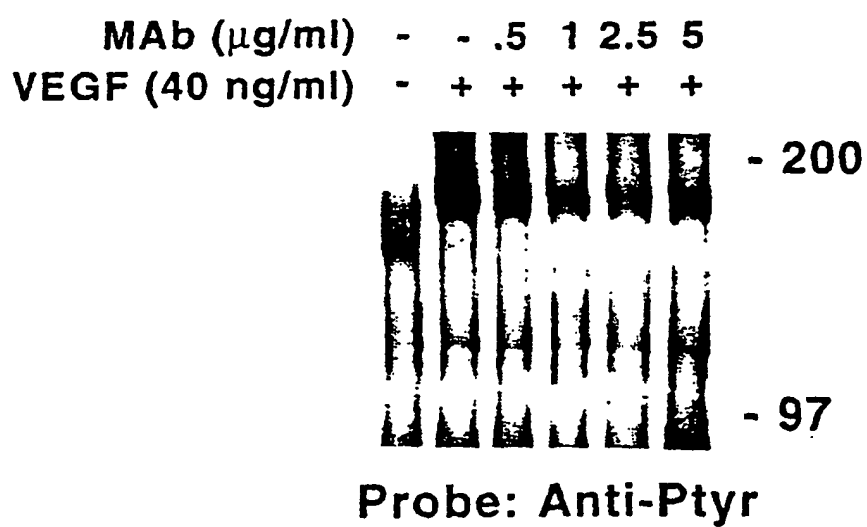
Western Blot of Flk-1/SEAPS Immunoprecipitation with MAb  
DC101



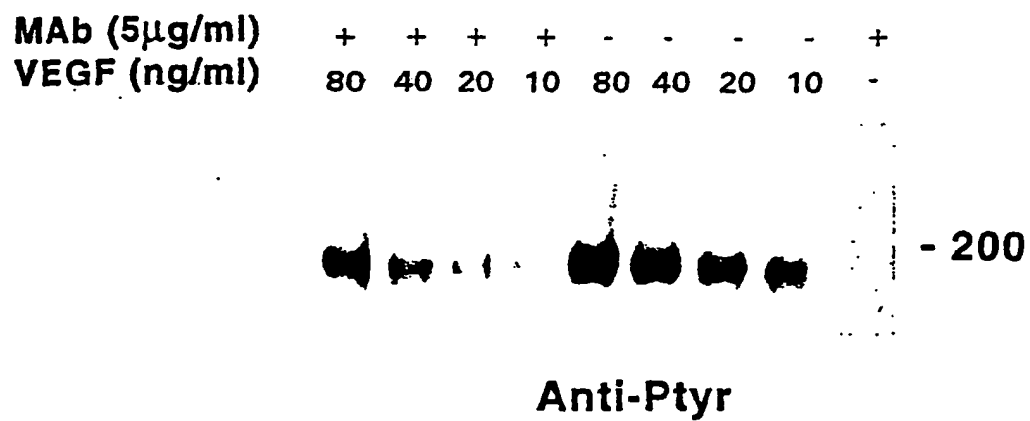
## FIGURE 2A

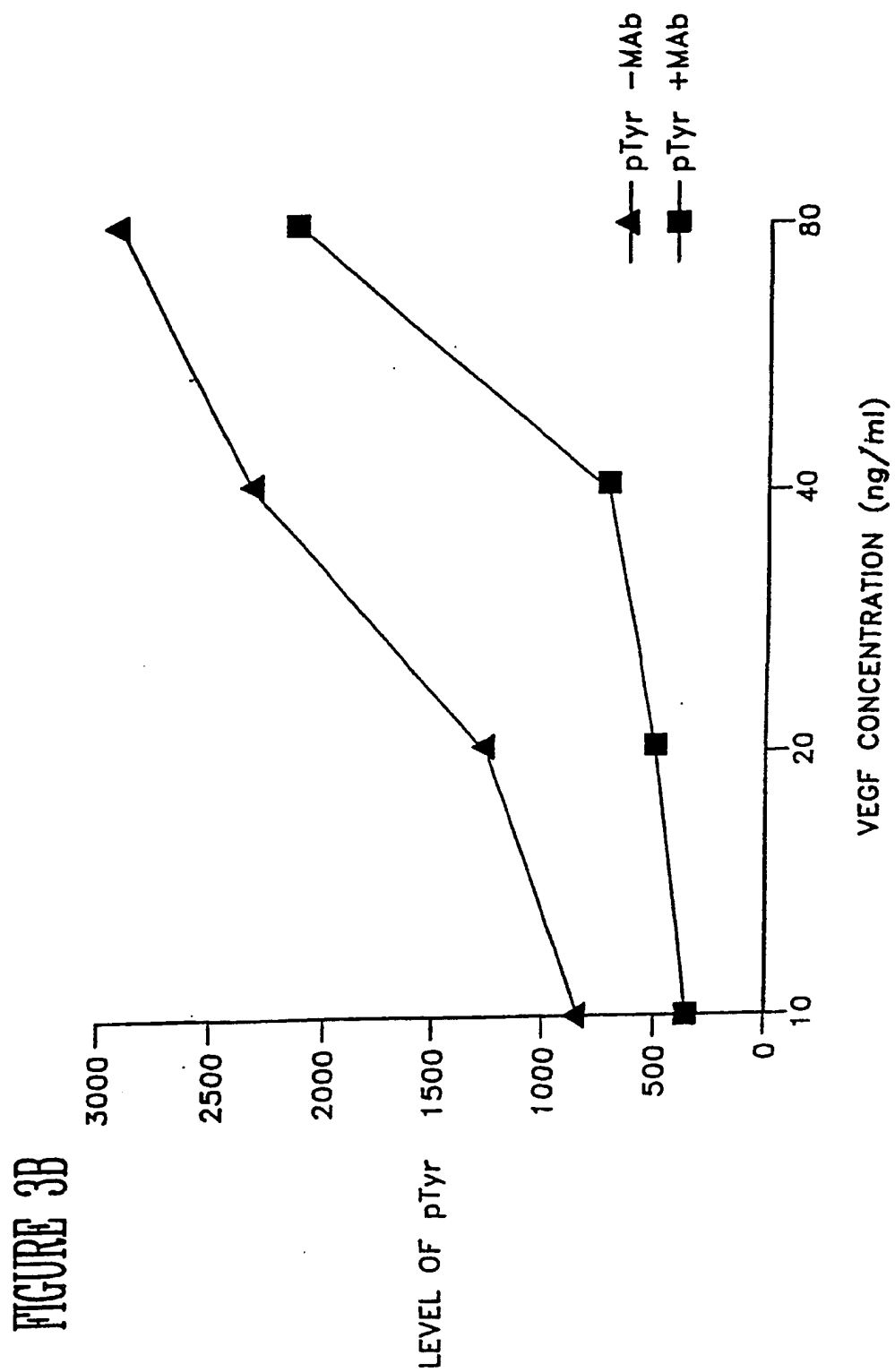


## FIGURE 2B



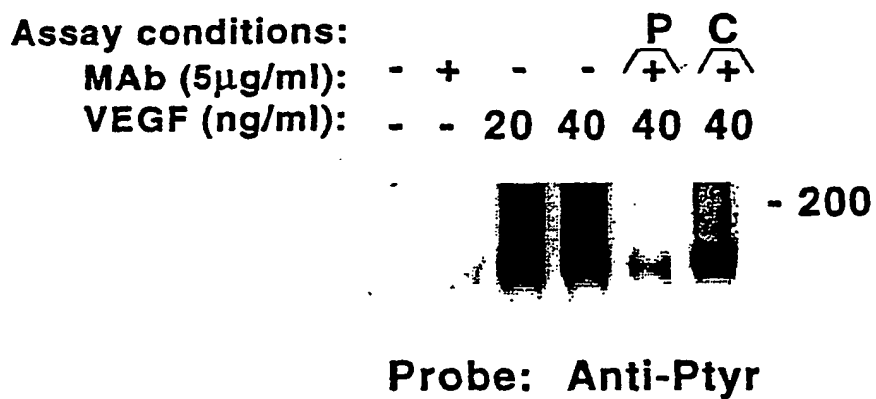
# FIGURE 3A





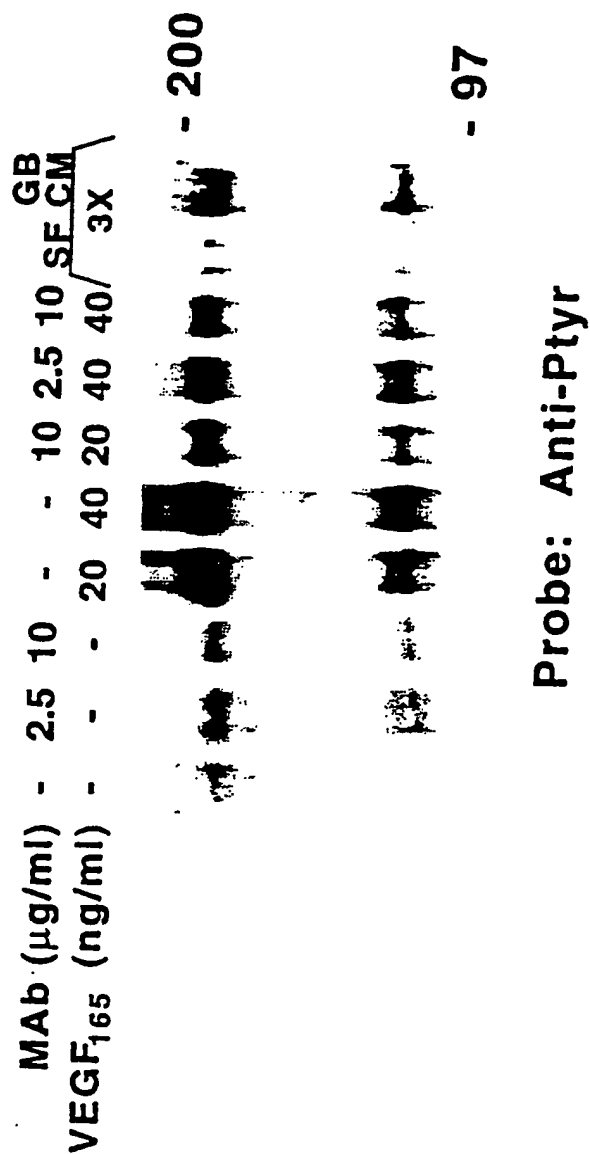
# FIGURE 4

Inhibition of VEGF-Flk-1/fms activation by pre-bound MAb  
DC101



Assay conditions: P: MAb pre-bound 15'; VEGF 15'  
C: Competitive assay; MAb  
+ VEGF 15'

## FIGURE 5



**FIGURE 6** FACS ANALYSIS OF Anti-flk-1 MAb BINDING TO flk-1/fms  
TRANSFECTED 3T3 CELLS (C441)

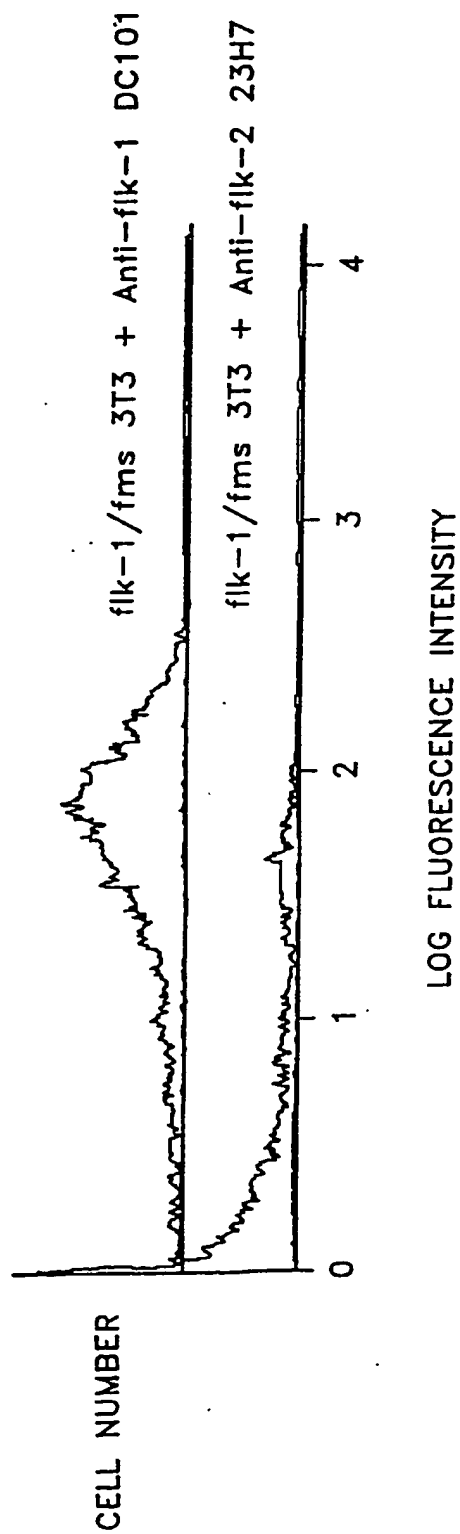
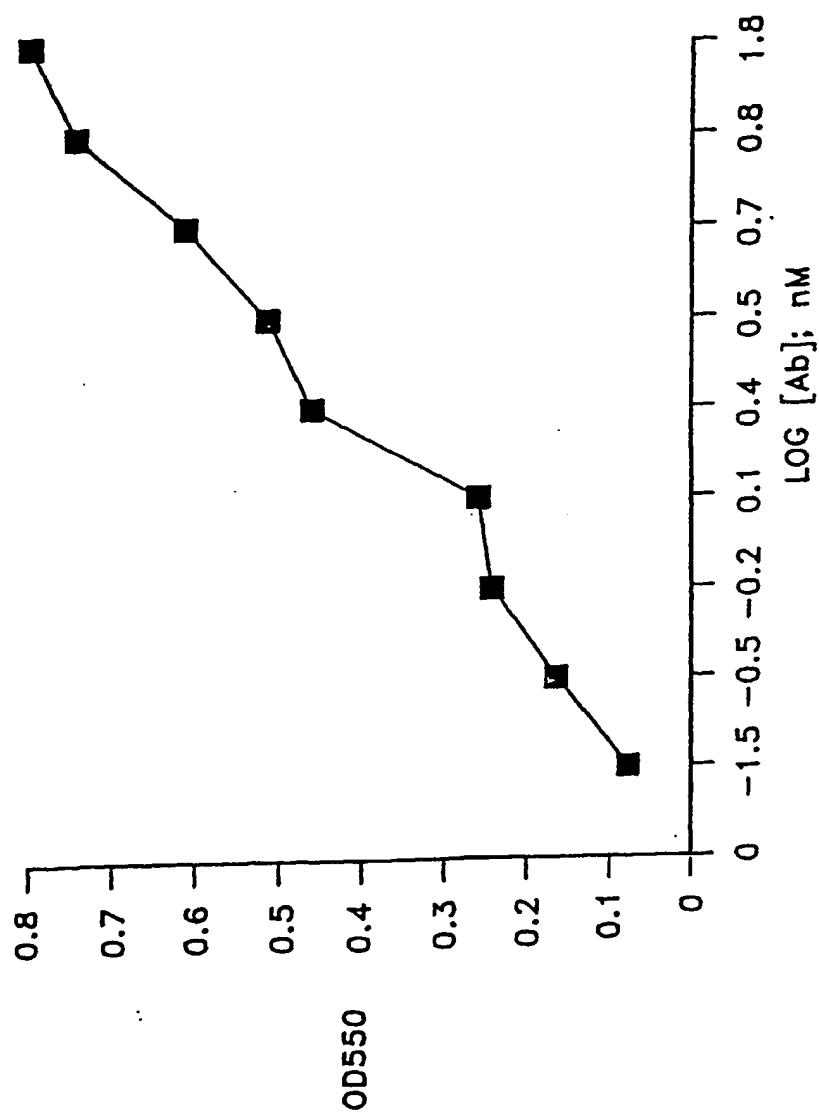


FIGURE 7 SATURATION BINDING OF MAb DC101 TO THE  
flk-1/fms RECEPTOR ON THE TRANSFECTED 3T3 CELL LINE C441



## FIGURE 8

Immunoprecipitation of phosphorylated Flk-1/fms from  
VEGF stimulated Flk-1/fms transfected 3T3 cells

**Antibody: 1 2 3 4**

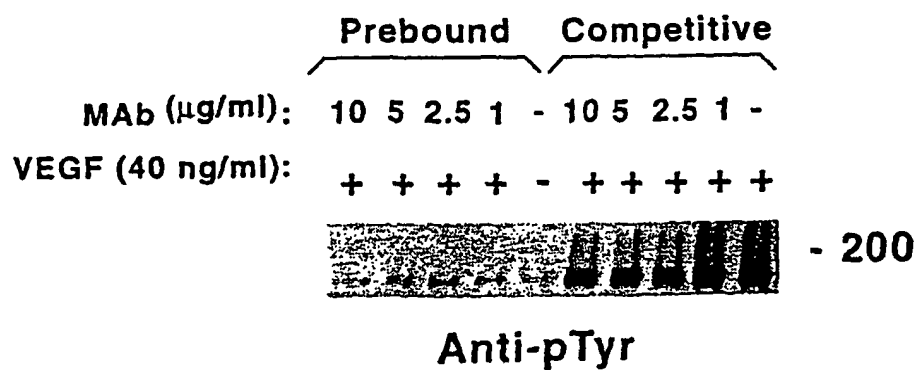
  - 200

**Anti-pTyr**

Antibodies: 1) Rat anti-flk-2 IgG<sub>2a</sub> 2A13  
2) Rat anti-flk-1 IgG<sub>1</sub> DC101  
3) Rat anti-flk-2 IgG<sub>1</sub> 23H7  
4) Rabbit anti-fms polyclonal IM 133

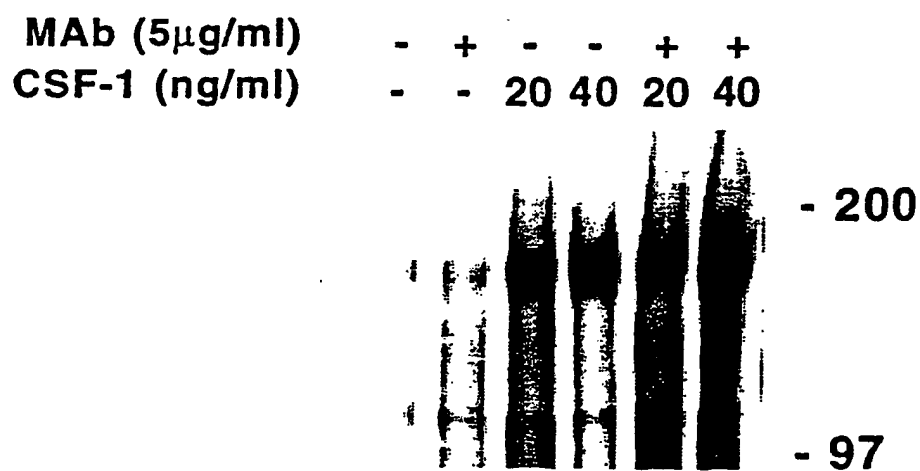
## FIGURE 9

Sensitivity of VEGF induced phosphorylation of the Flk-1/fms receptor to inhibition by MAb DC101



## FIGURE 10

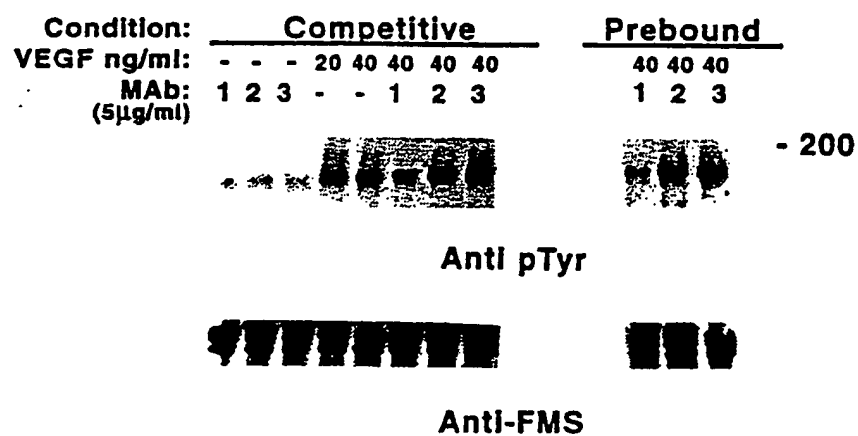
Effect of MAb DC101 on CSF-1 induced phosphorylation of the *fms* receptor



Probe: Anti-Ptyr

# FIGURE 11

Specificity of MAb DC101 neutralization of the activated Flk-1/fms receptor



Rat MAbs: 1) Anti-flk-1 IgG<sub>1</sub> DC101  
 2) Anti-flk-2 IgG<sub>1</sub> 23H7  
 3) Anti-flk-2 IgG<sub>2a</sub> 2A13

## FIGURE 12

Immunoprecipitation of phosphorylated receptor bands from  
VEGF stimulated HUVEC cells

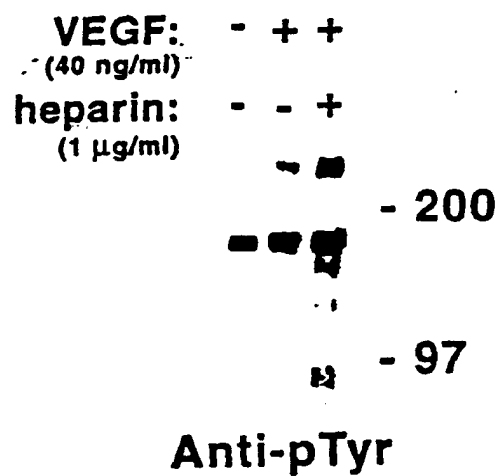
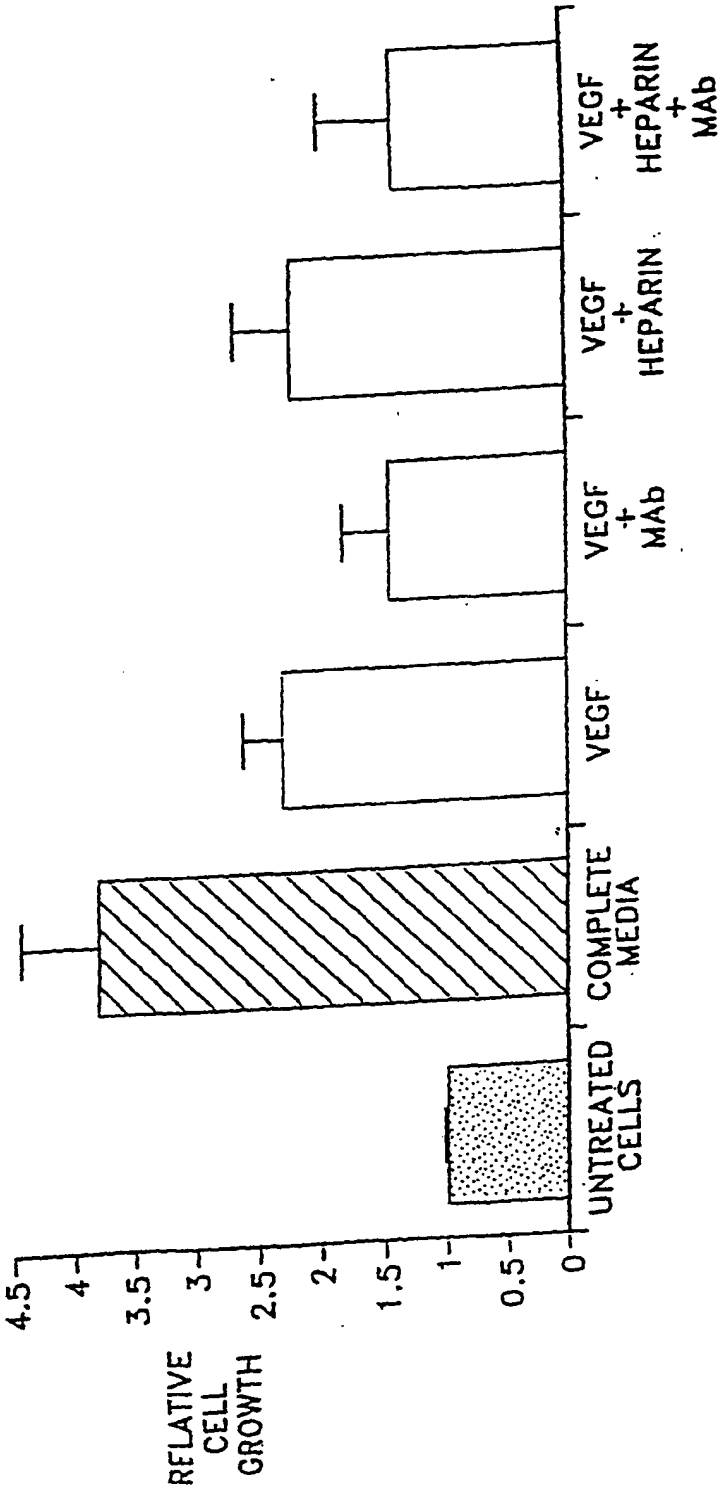


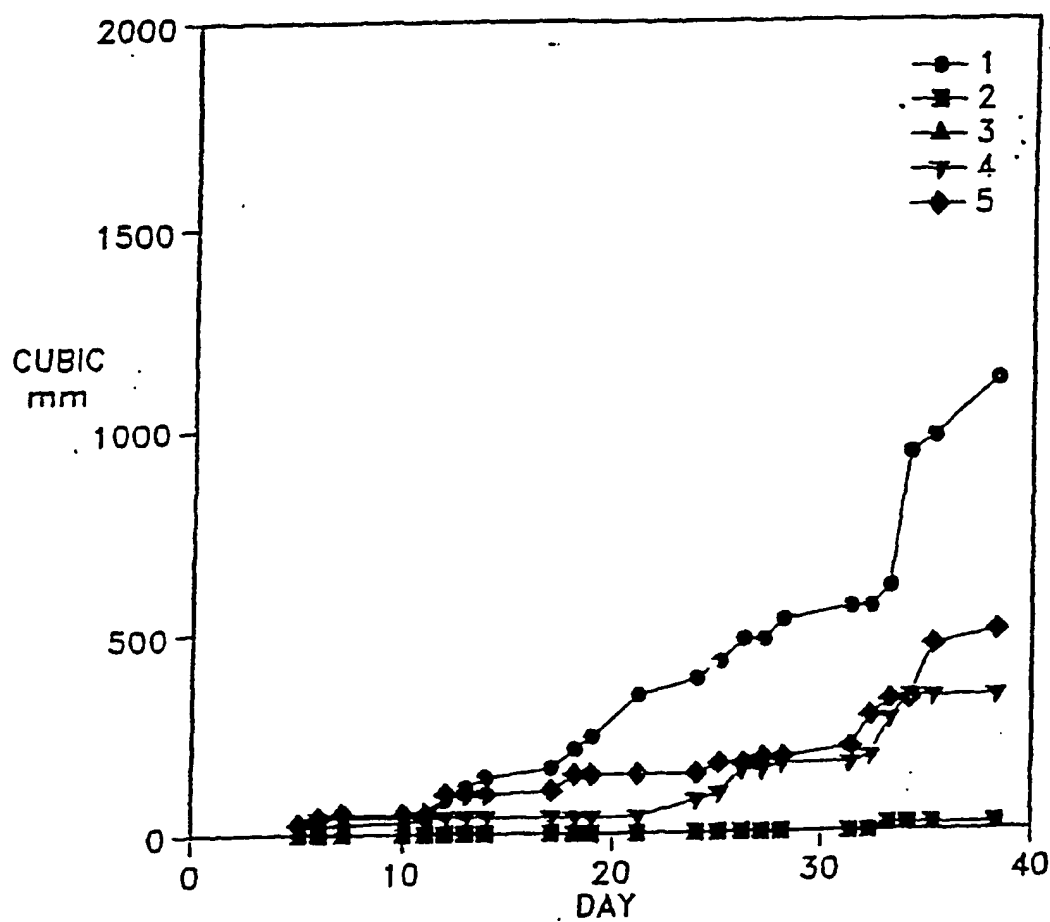
FIGURE 13

CONTROL 10% FBS+EC SUPPLEMENT VEGF



## FIGURE 14A

TREATMENT OF GLIOBLASTOMA  
XENOGRAPHS WITH RAT anti-flk-1 Mab



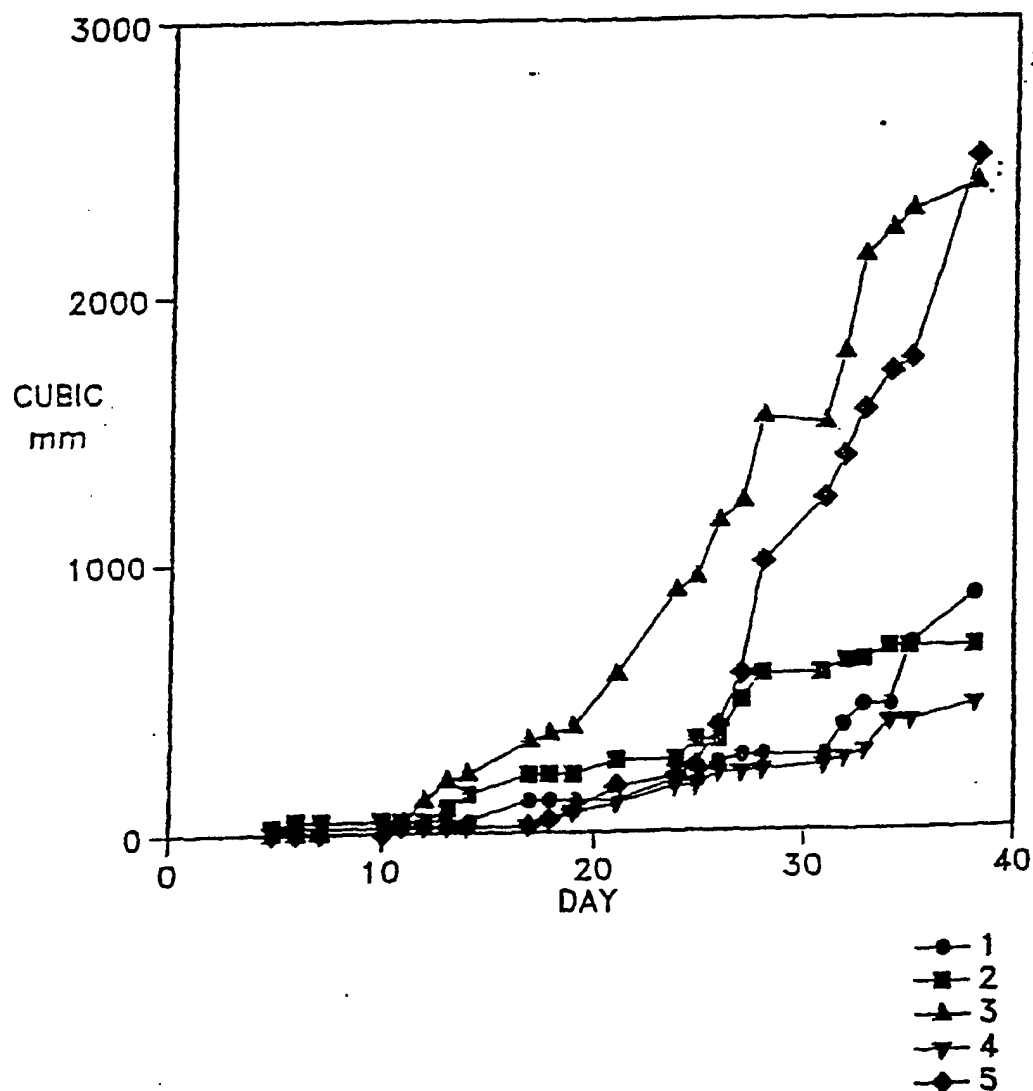
STATISTICAL ANALYSIS

flk-1 SLOPE = 16.09

p VALUE FOR flk-1 VERSUS flk-2 TUMOR SIZE = 0.0001

## FIGURE 14B

TREATMENT OF GLIOBLASTOMA  
XENOGRAFTS WITH RAT anti-flk-2 MAb

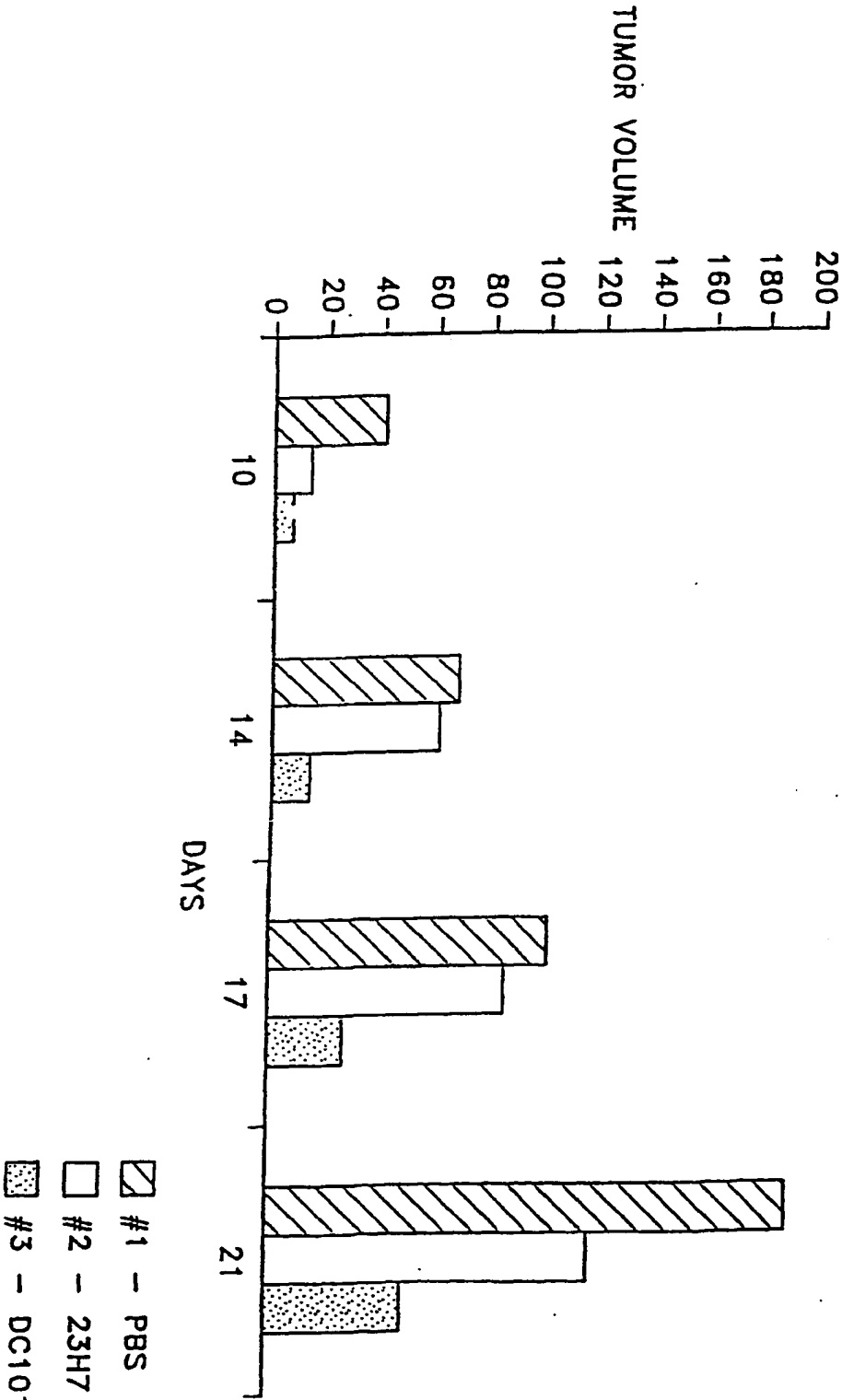


STATISTICAL ANALYSIS

flk-2 SLOPE = 37.39

P VALUE FOR flk-1 VERSUS flk-2 TUMOR SIZE = 0.0001

FIGURE 15



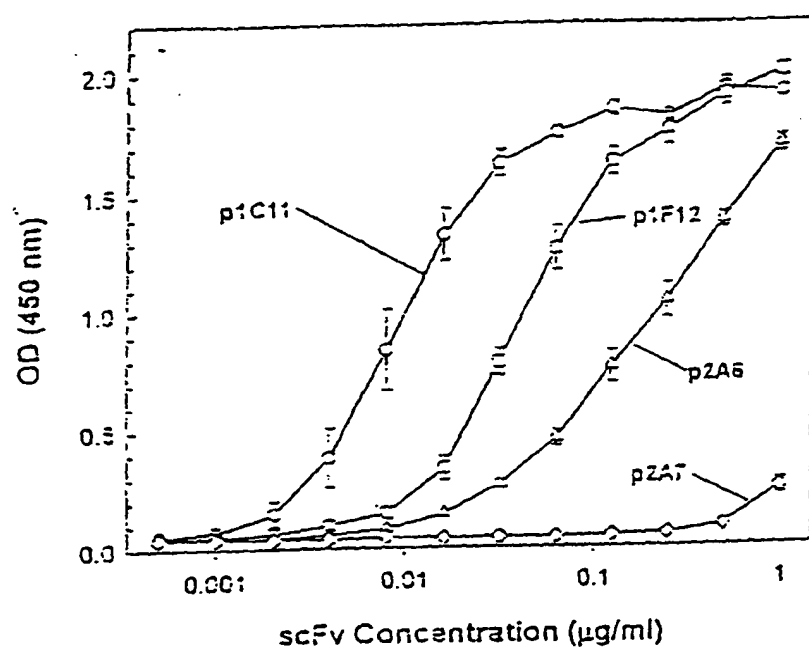


FIGURE 16

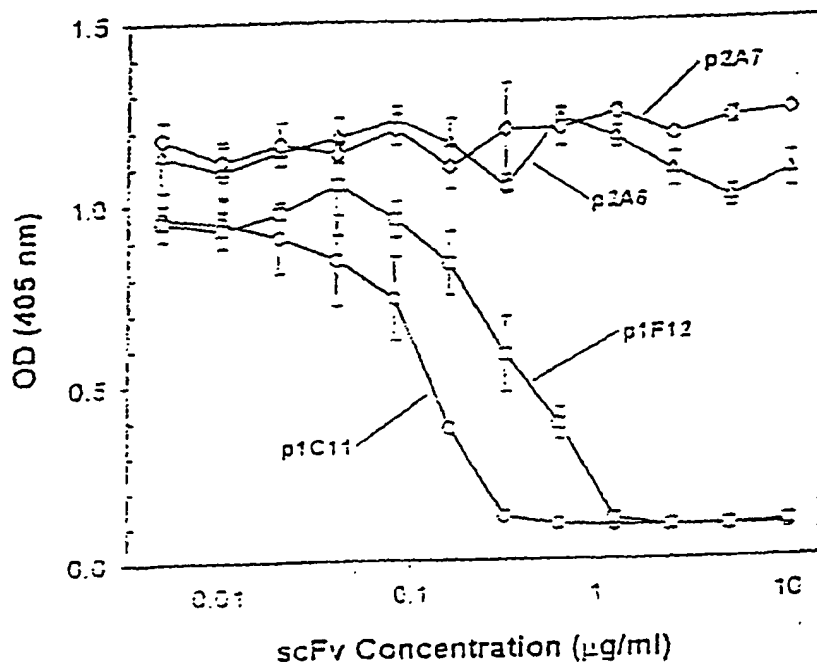


FIGURE 17

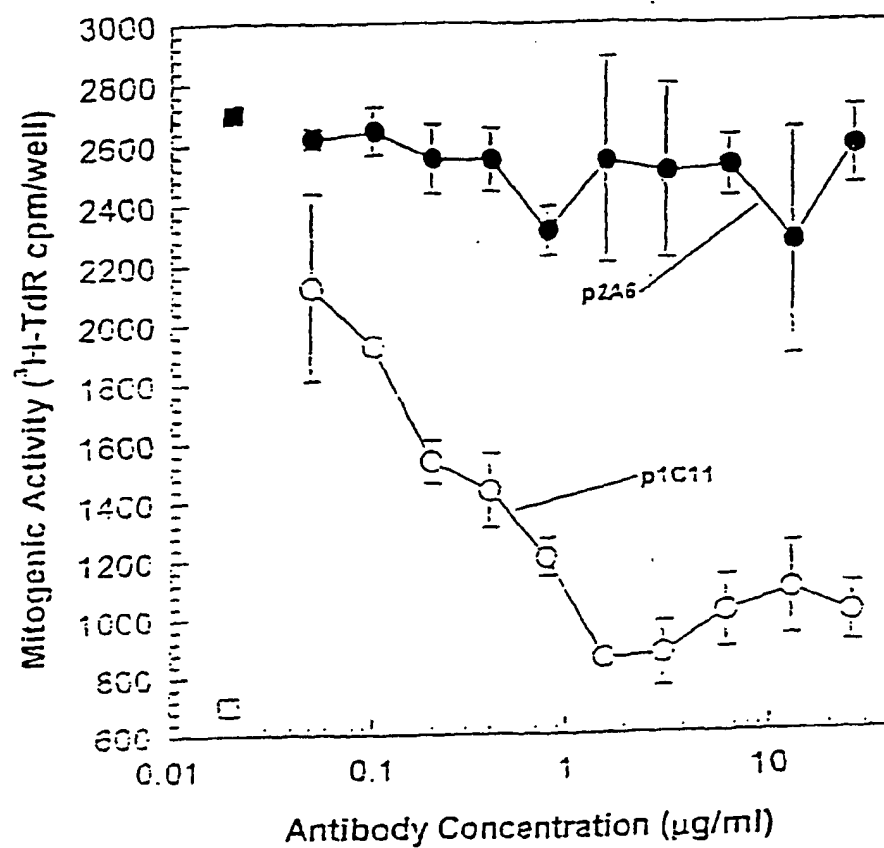


FIGURE 18

॥३॥

GGAACTCTGGGATGGTCTGGATCATGCTTTTCTAGTAGCACTGCACTGGAATACAT

M G W S C I I L E L V A T A T G V H

→ leader

TCTCAGGTCCTAGCTGCAACAGCTCTGGGGCAGAGCTTGTGGGGTCAGGGGCTCTAGTCAA

S ' Q V \_ X E Q Q S G A E L V G S G A S V X

五

[illegible]

U S C I S G E N Y F X D F Y M E W V K Q

63-2

[illegible]

2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20

CC-2-27

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Y A F K F G G K A T M T A B S S S N T A

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S E C R E T

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5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32

7-2-57

## 1.1.1.1

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22-00000

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— 2222 —

\_\_\_\_\_

**A B C D E F G H I J K L M N O P Q R S T U V**

10

11-11-76 10:00 AM 100-443887-1000

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١٠٠

\_\_\_\_\_

G C H U X L V I Z S T S Y E A U G V P N

**C23-22**

[illegible]

\_\_\_\_\_

F I C H E N O M M E : J A S T R O P H Y L A

C-3-21

3422

**၆၀၂၁၁၃ : ၆၄၇၅၈၉၁၂၃၄၅၆၇၈၉၀**

5 1 2 3 4 5 6

FIGURE 19.

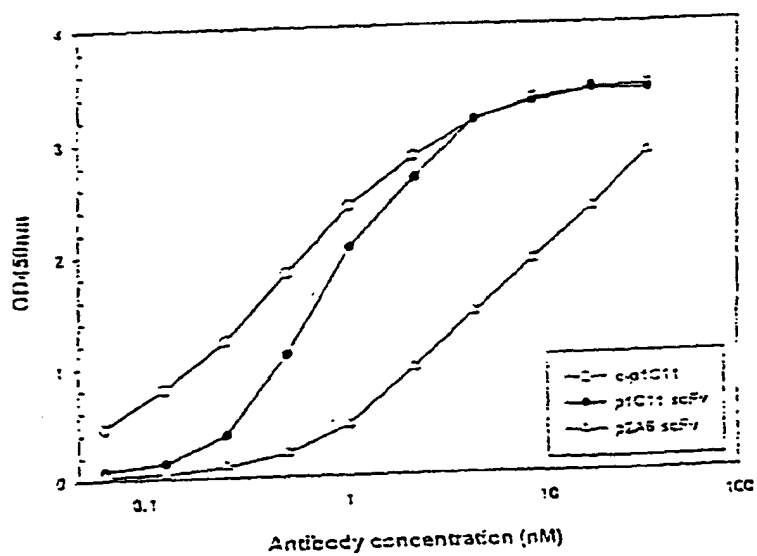


FIGURE 20

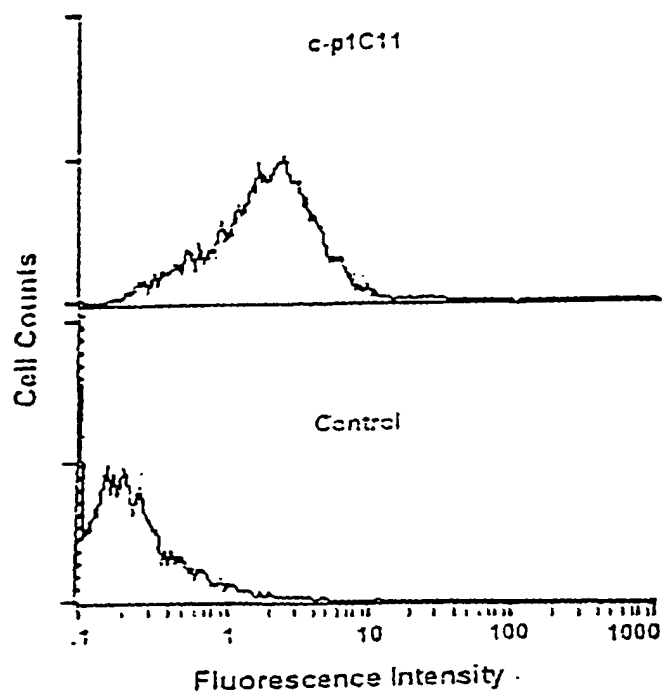


FIGURE 21

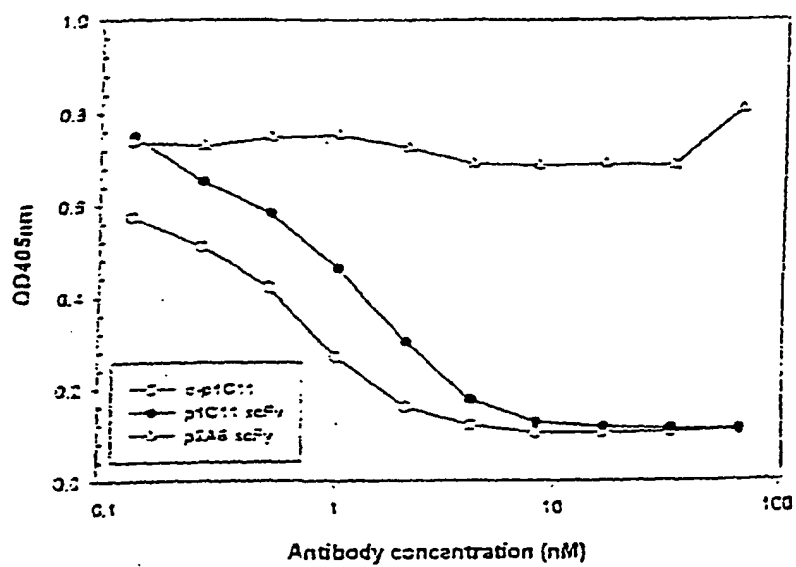


FIGURE 22

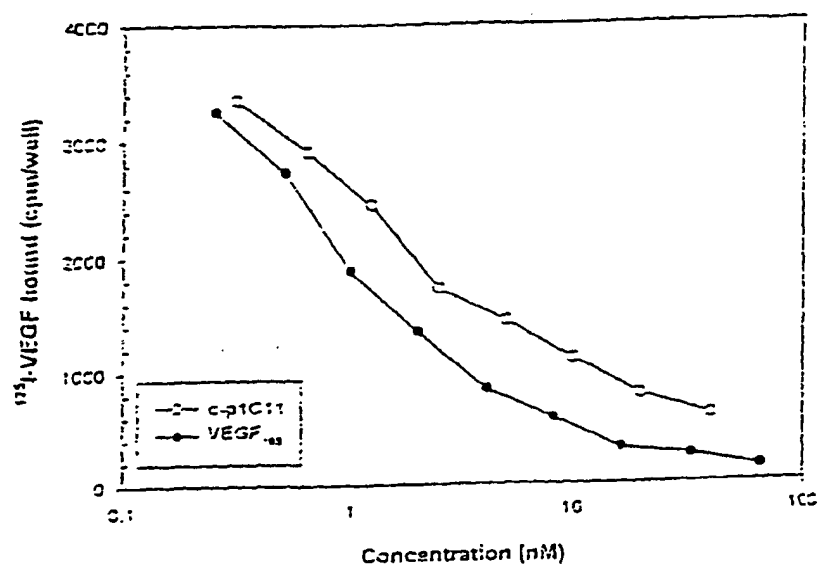


FIGURE 23

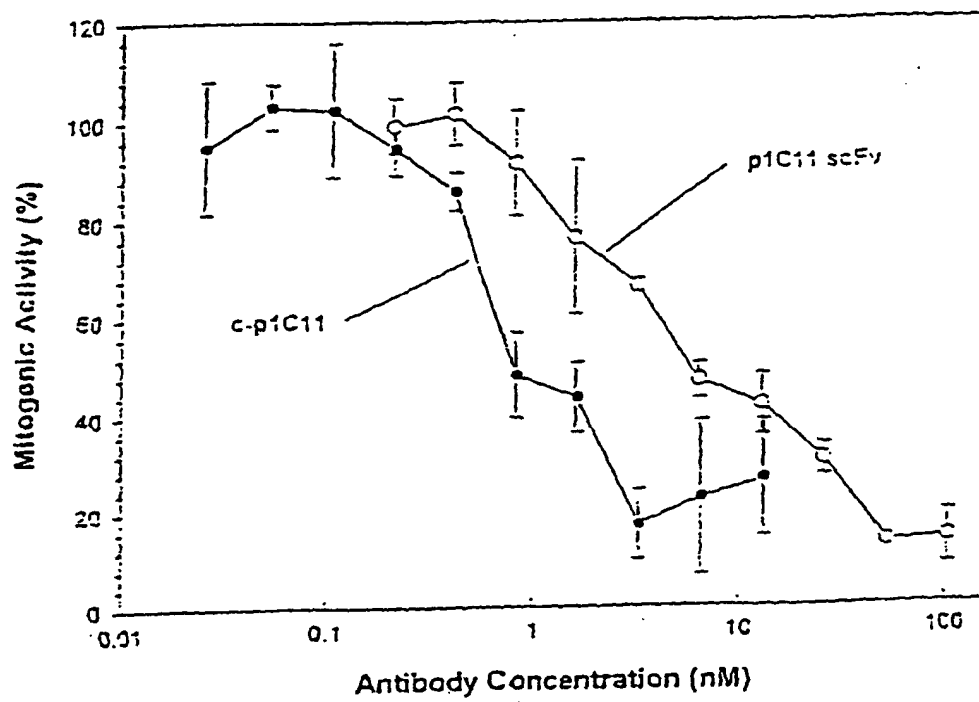


FIGURE 24

## SEQUENCE LISTING

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Goldstein, Neil I.  
<120> Combination Methods of Inhibiting Tumor Growth With a Vascular  
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<130> 11245/46276

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<211> 9

<212> PRT

<213> Human

<400> 3

Leu Gln His Asn Thr Phe Pro Pro Thr  
5

<210> 4

<211> 11

<212> PRT

<213> Human

<400> 4

Arg Ala Ser Gln Gly Ile Ser Ser Arg Leu Ala  
5 10

<210> 5

<211> 7

<212> PRT

<213> Human

&lt;400&gt; 5

Ala Ala Ser Ser Leu Gln Thr  
5

&lt;210&gt; 6

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 6

Gln Gln Ala Asn Arg Phe Pro Pro Thr  
5

&lt;210&gt; 7

&lt;211&gt; 14

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 7

Ala Gly Thr Thr Thr Asp Leu Thr Tyr Tyr Asp Leu Val Ser  
5 10

&lt;210&gt; 8

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 8

Asp Gly Asn Lys Arg Pro Ser  
5

&lt;210&gt; 9

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 9

Asn Ser Tyr Val Ser Ser Arg Phe Tyr Val  
5 10

&lt;210&gt; 10

&lt;211&gt; 13

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 10

Ser Gly Ser Thr Ser Asn Ile Gly Thr Asn Thr Ala Asn  
5 10

&lt;210&gt; 11

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 11

Asn Asn Asn Gln Arg Pro Ser  
5

&lt;210&gt; 12

&lt;211&gt; 12

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 12

Ala Ala Trp Asp Asp Ser Leu Asn Gly His Trp Val  
5 10

&lt;210&gt; 13

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 13

Gly Phe Thr Phe Ser Ser Tyr Ser Met Asn  
5 10

&lt;210&gt; 14

&lt;211&gt; 17

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 14

Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val Lys  
5 10 15

Gly  
17

&lt;210&gt; 15

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 15

Val Thr Asp Ala Phe Asp Ile  
5

&lt;210&gt; 16

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 16

Gly Gly Thr Phe Ser Ser Tyr Ala Ile Ser  
5 10



<210>	20
<211>	125
<212>	PRT
<213>	Human

<400> 20

Glu	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala
				5					10					15	
Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Gly	Thr	Phe	Ser	Ser	Tyr
			20					25					30		
Ala	Ile	Ser	Trp	Val	Arg	Gln	Ala	Pro	Gly	Gln	Gly	Leu	Glu	Trp	Met
		35					40					45			
Gly	Gly	Ile	Ile	Pro	Ile	Phe	Gly	Thr	Ala	Asn	Tyr	Ala	Gln	Lys	Phe
	50					55					60				
Gln	Gly	Arg	Val	Thr	Phe	Thr	Ala	Asp	Lys	Ser	Thr	Ser	Thr	Ala	Tyr
65					70					75					80
Met	Glu	Leu	Arg	Ser	Leu	Arg	Ser	Asp	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
				85					90					95	
Ala	Arg	Gly	Tyr	Asp	Tyr	Tyr	Asp	Ser	Ser	Gly	Val	Ala	Ser	Pro	Phe
			100					105					110		
Asp	Tyr	Trp	Gly	Gln	Gly	Thr	Leu	Val	Thr	Val	Ser	Ser			
		115					120					125			

<210>	21
<211>	333
<212>	DNA
<213>	Human

<400> 21

cag	tct	gtg	ctg	act	cag	cca	ccc	tca	gcg	tct	ggg	acc	ccc	ggg	cag	48
Gln	Ser	Val	Leu	Thr	Gln	Pro	Pro	Ser	Ala	Ser	Gly	Thr	Pro	Gly	Gln	
			5						10					15		
agg	gtc	acc	atc	tct	tgt	tct	gga	agc	acc	tcc	aac	atc	ggg	act	aat	96
Arg	Val	Thr	Ile	Ser	Cys	Ser	Gly	Ser	Thr	Ser	Asn	Ile	Gly	Thr	Asn	
			20					25					30			
act	gca	aac	tgg	ttc	cag	cag	ctc	cca	gga	acg	gcc	ccc	aaa	ctc	ctc	144
Thr	Ala	Asn	Trp	Phe	Gln	Gln	Leu	Pro	Gly	Thr	Ala	Pro	Lys	Leu	Leu	
		35					40					45				
atc	cac	aat	aat	aat	cag	cgg	ccc	tca	ggg	gtc	cct	gac	cga	ttc	tct	192
Ile	His	Asn	Asn	Asn	Gln	Arg	Pro	Ser	Gly	Val	Pro	Asp	Arg	Phe	Ser	
	50					55					60					
ggc	toc	aag	tct	ggc	acc	tca	gcc	tcc	ctg	gcc	atc	agt	ggg	ctc	cag	240
Gly	Ser	Lys	Ser	Gly	Thr	Ser	Ala	Ser	Leu	Ala	Ile	Ser	Gly	Leu	Gln	
65					70					75					80	
tct	gag	gat	gag	gct	gat	tat	tac	tgt	gca	gca	tgg	gat	gac	agg	ctg	288
Ser	Glu	Asp	Glu	Ala	Asp	Tyr	Tyr	Cys	Ala	Ala	Trp	Asp	Asp	Ser	Leu	
				85					90					95		

aat ggc cat tgg gtg ttc ggc gga ggg acc aag ctg acc gtc ctg 333  
 Asn Gly His Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu  
                   100                                  105                                  110

<210> 22  
 <211> 111  
 <212> PRT  
 <213> Human

<400> 22

Gln Ser Val Leu Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln  
                                   5                                  10                                  15  
 Arg Val Thr Ile Ser Cys Ser Gly Ser Thr Ser Asn Ile Gly Thr Asn  
                                   20                                  25                                  30  
 Thr Ala Asn Trp Phe Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu  
                                   35                                  40                                  45  
 Ile His Asn Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser  
                                   50                                  55                                  60  
 Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln  
 65                                  70                                  75                                  80  
 Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu  
                                   85                                  90                                  95  
 Asn Gly His Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu  
                                   100                                  105                                  110

<210> 23  
 <211> 348  
 <212> DNA  
 <213> Human

<400> 23

gag gtg cag ctg gtg cag tct ggg gga ggc ctg gtc aag cct ggg ggg 48  
 Glu Val Gln Leu Val Gln Ser Gly Gly Gly Leu Val Lys Pro Gly Gly  
                                   5                                  10                                  15  
 tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt agc tat 96  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr  
                                   20                                  25                                  30  
 agc atg aac tgg gtc cgc cag gct cca ggg aag ggg ctg gag tgg gtc 144  
 Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
                                   35                                  40                                  45  
 tca tcc att agt agt agt agt agt tac ata tac tac gca gac tca gtg 192  
 Ser Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val  
                                   50                                  55                                  60  
 aag ggc cga ttc acc atc tcc aga gac aac gcc aag aac tca ctg tat 240  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr  
 65                                  70                                  75                                  80  
 ctg caa atg aac agc ctg aga gcc gag gac acg gct gtg tat tac tgt 288  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys

	85	90	95	
gcg aga gtc aca gat gct ttt gat atc tgg ggc caa ggg aca atg gtc				336
Ala Arg Val Thr Asp Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val				
	100	105	110	
acc gtc tca agc				348
Thr Val Ser Ser				
	115			

<210> 24  
 <211> 116  
 <212> PRT  
 <213> Human

<400> 24

Glu Val Gln Leu Val Gln Ser Gly Gly Gly Leu Val Lys Pro Gly Gly	5	10	15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr	20	25	30
Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val	35	40	45
Ser Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val	50	55	60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr	65	70	75
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys	85	90	95
Ala Arg Val Thr Asp Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val	100	105	110
Thr Val Ser Ser	115		

<210> 25  
 <211> 321  
 <212> DNA  
 <213> Human

<400> 25

gaa att gtg atg aca cag tct cca gcc acc ctg tct ttg tct cca ggg				48
Glu Ile Val Met Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser Pro Gly				
	5	10	15	
gaa aga gcc acc ctc tcc tgc agg gcc agt cag agt gtt agc agc tac				96
Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Tyr				
	20	25	30	
tta gcc tgg tac caa cag aaa cct gcc cag gct ccc agg ctc ctc atc				144
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile				
	35	40	45	
tat gat tca tcc aac agg gcc act gcc atc cca gcc aga ttc agt ggc				192
Tyr Asp Ser Ser Asn Arg Ala Thr Gly Ile Pro Ala Arg Phe Ser Gly				

50	55	60	
agt ggg tct ggg aca gac ttc act ctc acc atc agc agc cta gag cct			240
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro			
65	70	75	80
gaa gat ttt gca act tat tac tgt cta cag cat aac act ttt cct ccg			288
Glu Asp Phe Ala Thr Tyr Tyr Cys Leu Gln His Asn Thr Phe Pro Pro			
	85	90	95
acg ttc ggc caa ggg acc aag gtg gaa atc aaa			321
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys			
	100	105	

<210> 26  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 26

Glu Ile Val Met Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser Pro Gly			
	5	10	15
Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Tyr			
	20	25	30
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile			
	35	40	45
Tyr Asp Ser Ser Asn Arg Ala Thr Gly Ile Pro Ala Arg Phe Ser Gly			
	50	55	60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro			
	65	70	75
Glu Asp Phe Ala Thr Tyr Tyr Cys Leu Gln His Asn Thr Phe Pro Pro			
	85	90	95
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys			
	100	105	

<210> 27  
 <211> 348  
 <212> DNA  
 <213> Human

<400> 27

gag gtc cag ctg gtg cag tct ggg gga ggc ctg gtc aag cct ggg ggg			48
Glu Val Gln Leu Val Gln Ser Gly Gly Gly Leu Val Lys Pro Gly Gly			
	5	10	15
tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt agc tat			96
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr			
	20	25	30
agc atg aac tgg gtc cgc cag gct cca ggg aag ggg ctg gag tgg gtc			144
Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val			
	35	40	45
tca tcc att agt agt agt agt agt tac ata tac tac gca gac tca gtg			192

Ser Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val  
     50                                    55                                    60

aag ggc cga ttc acc atc tcc aga gac aac gcc aag aac tca ctg tat 240  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr  
     65                                    70                                    75                                    80

ctg caa atg aac agc ctg aga gcc gag gac acg gct gtg tat tac tgt 288  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
                                     85                                    90                                    95

gcg aga gtc aca gat gct ttt gat atc tgg ggc caa ggg aca atg gtc 336  
 Ala Arg Val Thr Asp Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val  
                                     100                                    105                                    110

acc gtc tca agc 348  
 Thr Val Ser Ser  
     115

<210> 28  
 <211> 330  
 <212> DNA  
 <213> Human

<400> 28

cag tct gcc ctg act cag cct gcc tcc ctg tct ggg tct cct gga cag 48  
 Gln Ser Ala Leu Thr Gln Pro Ala Ser Leu Ser Gly Ser Pro Gly Gln  
                                     5                                    10                                    15

tcg atc acc atc tcc tgc gct gga acc acc act gat ctt aca tat tat 96  
 Ser Ile Thr Ile Ser Cys Ala Gly Thr Thr Thr Asp Leu Thr Tyr Tyr  
                                     20                                    25                                    30

gac ctt gtc tcc tgg tac caa cag cac cca ggc caa gca ccc aaa ctc 144  
 Asp Leu Val Ser Trp Tyr Gln Gln His Pro Gly Gln Ala Pro Lys Leu  
                                     35                                    40                                    45

gtg att tat gac ggc aat aag cgg ccc tca gga gtt tct aat cgc ttc 192  
 Val Ile Tyr Asp Gly Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe  
     50                                    55                                    60

tct ggc tcc aag tct ggc aac acg gcc tcc ctg aca atc tct gga ctc 240  
 Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu  
     65                                    70                                    75                                    80

cag gct gag gac gag gct gat tat tac tgc aac tca tat gta agc agc 288  
 Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Asn Ser Tyr Val Ser Ser  
                                     85                                    90                                    95

agg ttt tat gtc ttc gga act ggg acc aag gtc acc gtc cta 330  
 Arg Phe Tyr Val Phe Gly Thr Gly Thr Lys Val Thr Val Leu  
                                     100                                    105                                    110

<210> 29  
 <211> 110  
 <212> PRT  
 <213> Human

<400> 29

Gln	Ser	Ala	Leu	Thr	Gln	Pro	Ala	Ser	Leu	Ser	Gly	Ser	Pro	Gly	Gln
				5					10					15	
Ser	Ile	Thr	Ile	Ser	Cys	Ala	Gly	Thr	Thr	Thr	Asp	Leu	Thr	Tyr	Tyr
			20					25					30		
Asp	Leu	Val	Ser	Trp	Tyr	Gln	Gln	His	Pro	Gly	Gln	Ala	Pro	Lys	Leu
		35					40					45			
Val	Ile	Tyr	Asp	Gly	Asn	Lys	Arg	Pro	Ser	Gly	Val	Ser	Asn	Arg	Phe
	50					55					60				
Ser	Gly	Ser	Lys	Ser	Gly	Asn	Thr	Ala	Ser	Leu	Thr	Ile	Ser	Gly	Leu
65					70					75					80
Gln	Ala	Glu	Asp	Glu	Ala	Asp	Tyr	Tyr	Cys	Asn	Ser	Tyr	Val	Ser	Ser
				85					90					95	
Arg	Phe	Tyr	Val	Phe	Gly	Thr	Gly	Thr	Lys	Val	Thr	Val	Leu		
			100					105					110		

<210>	30
<211>	348
<212>	DNA
<213>	Human

<400> 30

[illegible]

<210> 31  
<211> 116

<212> PRT  
<213> Human

<400> 31

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Glu Val Gln Leu Val Gln Ser Gly Gly Gly Leu Val Lys Pro Gly Gly
      5              10              15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
      20              25              30
Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
      35              40              45
Ser Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val
      50              55              60
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asp Ser Leu Tyr
      65              70              75
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
      85              90              95
Ala Arg Val Thr Asp Ala Phe Asp Ile Trp Gly Gln Gly Thr Met Val
      100             105             110
Thr Val Ser Ser
      115

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<210> 32  
<211> 321  
<212> DNA  
<213> Human

<400> 32

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gac atc cag ttg acc cag tct cca tct tct gtg tct gca tct gta gga      48
Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
      5              10              15

gac aga gtc acc atc act tgt cgg gcg agt cag ggt att agt agt cgg      96
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser Arg
      20              25              30

tta gcc tgg tat cag cag aaa cca ggg aaa gcc cct aag ctc ctg atc      144
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
      35              40              45

tat gct gca tcc agt ttg caa act ggg gtc cca tca agg ttc agc ggc      192
Tyr Ala Ala Ser Ser Leu Gln Thr Gly Val Pro Ser Arg Phe Ser Gly
      50              55              60

agt gga tct ggg aca gat ttc act ctc act atc agc agc ctg cag cct      240
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
      65              70              75

gaa gat ttt gca act tac tat tgt caa cag gct aac agg ttc cct ccg      288
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ala Asn Arg Phe Pro Pro
      85              90              95

act ttc ggc cct ggg acc aaa gtg gat atc aaa      321
Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
      100             105

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<210> 33  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 33

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Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
                    5                      10                      15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser Arg
                20                      25                      30

Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                35                      40                      45

Tyr Ala Ala Ser Ser Leu Gln Thr Gly Val Pro Ser Arg Phe Ser Gly
                50                      55                      60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
        65                      70                      75                      80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ala Asn Arg Phe Pro Pro
                85                      90                      95

Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
                100                      105
  
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<210> 34  
 <211> 333  
 <212> DNA  
 <213> Human

<400> 34

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cag tct gtc gtg acg cag ccg ccc tca gtg tct ggg gcc cca ggg cag      48
Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
                    5                      10                      15

agg gtc acc atc tcc tgc act ggg agc cac tcc aac ttc ggg gca gga      96
Arg Val Thr Ile Ser Cys Thr Gly Ser His Ser Asn Phe Gly Ala Gly
                20                      25                      30

act gat gta cat tgg tac caa cac ctt cca gga aca gcc ccc aga ctc     144
Thr Asp Val His Trp Tyr Gln His Leu Pro Gly Thr Ala Pro Arg Leu
                35                      40                      45

ctc att cat gga gac agt aat cgg ccc tcc ggg gtc cct gac cga ttc     192
Leu Ile His Gly Asp Ser Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
                50                      55                      60

tct ggc tcc agg tct ggc acc tca gcc tcc ctg gcc atc act ggg ctc     240
Ser Gly Ser Arg Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu
                65                      70                      75                      80

cgg gtt gag gat gag gct gat tat tac tgt cag tcg tat gac tat ggc     288
Arg Val Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Gly
                85                      90                      95

ctg aga ggt tgg gtg ttc ggc ggc ggg acc aag ctg acc gtc ctt      333
Leu Arg Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
                100                      105                      110
  
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<210> 35  
 <211> 111  
 <212> PRT  
 <213> Human

<400> 35

Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln  
                                   5                                  10                                  15  
 Arg Val Thr Ile Ser Cys Thr Gly Ser His Ser Asn Phe Gly Ala Gly  
                                   20                                  25                                  30  
 Thr Asp Val His Trp Tyr Gln His Leu Pro Gly Thr Ala Pro Arg Leu  
                                   35                                  40                                  45  
 Leu Ile His Gly Asp Ser Asn Arg Pro Ser Gly Val Pro Asp Arg Phe  
                                   50                                  55                                  60  
 Ser Gly Ser Arg Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu  
                                   65                                  70                                  75                                  80  
 Arg Val Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Gly  
                                   85                                  90                                  95  
 Leu Arg Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu  
                                   100                                  105                                  110

<210> 36  
 <211> 321  
 <212> DNA  
 <213> Human

<400> 36

gat gtt gtg atg act cag tct cca tcg tcc ctg tct gca tct gta ggg 48  
 Asp Val Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
                                   5                                  10                                  15  
 gac aga gtc acc atc act tgc cgg gca agt cag aac att aac aac tat 96  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Asn Ile Asn Asn Tyr  
                                   20                                  25                                  30  
 tta aat tgg tat caa cag aaa cca gga aaa gcc cct aag ctc ctg atc 144  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
                                   35                                  40                                  45  
 tat gct gcc tcc act ttg caa agt ggg gtc cca tca agg ttc agt ggc 192  
 Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
                                   50                                  55                                  60  
 agt gga tct ggg aca gat ttc act ctc acc atc acc agc cta cag cct 240  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Thr Ser Leu Gln Pro  
                                   65                                  70                                  75                                  80  
 gaa gat tct gca act tat tac tgc caa cag tat tcc cgt tat cct ccc 288  
 Glu Asp Ser Ala Thr Tyr Tyr Cys Gln Gln Tyr Ser Arg Tyr Pro Pro  
                                   85                                  90                                  95  
 act ttc ggc gga ggg acc aag gtg gag atc aca 321  
 Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Thr  
                                   100                                  105

<210> 37  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 37

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Asp Val Val Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
              5                      10                      15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Asn Ile Asn Asn Tyr
              20                      25                      30

Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
              35                      40                      45

Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
              50                      55                      60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Thr Ser Leu Gln Pro
              65                      70                      75                      80

Glu Asp Ser Ala Thr Tyr Tyr Cys Gln Gln Tyr Ser Arg Tyr Pro Pro
              85                      90                      95

Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Thr
              100                      105
  
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<210> 38  
 <211> 330  
 <212> DNA  
 <213> Human

<400> 38

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cag tct gcc ctg act cag cct gcc tcc gtg tct ggg tct cgt gga cag      48
Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Arg Gly Gln
              5                      10                      15

tcg atc acc ctc tcc tgc acc ggc tcc agc act gat gtg ggt aat tat      96
Ser Ile Thr Leu Ser Cys Thr Gly Ser Ser Thr Asp Val Gly Asn Tyr
              20                      25                      30

aac tat atc tcc tgg tac caa caa cac cca ggc caa gcc ccc aaa ctc     144
Asn Tyr Ile Ser Trp Tyr Gln Gln His Pro Gly Gln Ala Pro Lys Leu
              35                      40                      45

ttg att tac gat gtc act agt cgg ccc tca ggt gtt tct gat cgc ttc     192
Leu Ile Tyr Asp Val Thr Ser Arg Pro Ser Gly Val Ser Asp Arg Phe
              50                      55                      60

tct ggc tcc aag tca ggc ctc acg gcc tcc ctg acc atc tct gga ctc     240
Ser Gly Ser Lys Ser Gly Leu Thr Ala Ser Leu Thr Ile Ser Gly Leu
              65                      70                      75                      80

cag cct gaa gac gag gct gac tat tac tgc aac tcc tat tct gcc acc     288
Gln Pro Glu Asp Glu Ala Asp Tyr Tyr Cys Asn Ser Tyr Ser Ala Thr
              85                      90                      95

gac act ctt gtt ttt ggc gga ggg acc aag ctg acc gtc cta             330
Asp Thr Leu Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
              100                      105                      110
  
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<210> 39  
 <211> 110  
 <212> PRT  
 <213> Human

<400> 39

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Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Arg Gly Gln
      5              10              15

Ser Ile Thr Leu Ser Cys Thr Gly Ser Ser Thr Asp Val Gly Asn Tyr
      20              25              30

Asn Tyr Ile Ser Trp Tyr Gln Gln His Pro Gly Gln Ala Pro Lys Leu
      35              40              45

Leu Ile Tyr Asp Val Thr Ser Arg Pro Ser Gly Val Ser Asp Arg Phe
      50              55              60

Ser Gly Ser Lys Ser Gly Leu Thr Ala Ser Leu Thr Ile Ser Gly Leu
      65              70              75              80

Gln Pro Glu Asp Glu Ala Asp Tyr Tyr Cys Asn Ser Tyr Ser Ala Thr
      85              90              95

Asp Thr Leu Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
      100              105              110
  
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<210> 40  
 <211> 333  
 <212> DNA  
 <213> Human

<400> 40

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cag gct gtg ctg act cag ccg tcc tca gtg tct ggg gcc cca gga cag      48
Gln Ala Val Leu Thr Gln Pro Ser Ser Val Ser Gly Ala Pro Gly Gln
      5              10              15

agg gtc acc atc tcc tgc act ggg caa agc tcc aat atc ggg gca gat      96
Arg Val Thr Ile Ser Cys Thr Gly Gln Ser Ser Asn Ile Gly Ala Asp
      20              25              30

tat gat gta cat tgg tac cag caa ttt cca gga aca gcc ccc aaa ctc      144
Tyr Asp Val His Trp Tyr Gln Gln Phe Pro Gly Thr Ala Pro Lys Leu
      35              40              45

ctc atc tat ggt cac aac aat cgg ccc tca ggg gtc cct gac cga ttc      192
Leu Ile Tyr Gly His Asn Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
      50              55              60

tct ggc tcc aag tct ggc acc tca gtc tcc ctg gtc atc agt ggg ctc      240
Ser Gly Ser Lys Ser Gly Thr Ser Val Ser Leu Val Ile Ser Gly Leu
      65              70              75              80

cag gct gag gat gag gct gat tat tat tgc cag tcc tat gac agc agt      288
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Ser Ser
      85              90              95

cta agt ggt ttg gta ttc ggc gga ggg acc aag gtg acc gtc cta      333
Leu Ser Gly Leu Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu
      100              105              110
  
```



<210> 43  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 43

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Asp Ile Gln Leu Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
                    5                      10              15

Asp Ser Val Thr Ile Thr Cys Arg Ala Ser Gln Asp Ile Ser Ser Trp
                20                      25              30

Leu Ala Trp Tyr Gln Gln Lys Pro Gly Glu Ala Pro Lys Leu Leu Ile
                35                      40              45

Tyr Ala Ala Ser Leu Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
                50                      55              60

Ser Gly Ser Gly Thr Asp Phe Ala Leu Thr Ile Asn Ser Leu Gln Pro
                65                      70              75              80

Glu Asp Phe Ala Thr Tyr Phe Cys Gln Gln Ala Asp Ser Phe Pro Pro
                85                      90              95

Thr Phe Gly Gln Gly Thr Arg Leu Glu Ile Lys
                100                      105
  
```

<210> 44  
 <211> 321  
 <212> DNA  
 <213> Human

<400> 44

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gac atc gag ttg acc cag tct cca tct tcc gtg tct gca tct gtg gga      48
Asp Ile Glu Leu Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
                    5                      10              15

gac aga gtc acc ctc act tgt cgg gcg agt cag agt att aag agg tgg      96
Asp Arg Val Thr Leu Thr Cys Arg Ala Ser Gln Ser Ile Lys Arg Trp
                20                      25              30

tta gcc tgg tat cag cag aaa cca ggg aag gcc cct agg ctc ctc atc     144
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Arg Leu Leu Ile
                35                      40              45

tat gct gca tcc act ttg caa agt ggg gtc cca tca agg ttc agc ggc     192
Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
                50                      55              60

ggg gga tct ggg aca gat ttc act ctc acc atc aac agc ctg cag cct     240
Gly Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Asn Ser Leu Gln Pro
                65                      70              75              80

gaa gat ttt gca att tac tac tgt caa cag gct aac agt ttc cct ccc     288
Glu Asp Phe Ala Ile Tyr Tyr Cys Gln Gln Ala Asn Ser Phe Pro Pro
                85                      90              95

act ttc ggc cct ggg acc aaa gtg gat atc aaa                        321
Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
                100                      105
  
```

<210> 45  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 45

```

Asp Ile Glu Leu Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
              5                      10                      15

Asp Arg Val Thr Leu Thr Cys Arg Ala Ser Gln Ser Ile Lys Arg Trp
              20                      25                      30

Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Arg Leu Leu Ile
              35                      40                      45

Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
              50                      55                      60

Gly Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Asn Ser Leu Gln Pro
              65                      70                      75                      80

Glu Asp Phe Ala Ile Tyr Tyr Cys Gln Gln Ala Asn Ser Phe Pro Pro
              85                      90                      95

Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
              100                      105

```

<210> 46  
 <211> 333  
 <212> DNA  
 <213> Human

<400> 46

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cag tct gtc gtg acg cag ccg ccc tca gtg tct ggg gcc cca ggg cag      48
Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
              5                      10                      15

agg gtc acc atc tcc tgc agt ggg agc agg tcc aac atc ggg gca cac      96
Arg Val Thr Ile Ser Cys Ser Gly Ser Arg Ser Asn Ile Gly Ala His
              20                      25                      30

tat gaa gtc cag tgg tac cag cag ttt ccg gga gca gcc ccc aaa ctc      144
Tyr Glu Val Gln Trp Tyr Gln Phe Pro Gly Ala Ala Pro Lys Leu
              35                      40                      45

ctc atc tat ggt gac acc aat cgg ccc tca ggg gtc cct gac cga ttc      192
Leu Ile Tyr Gly Asp Thr Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
              50                      55                      60

tct gcc tcc cac tct ggc acc tca gcc tcc ctt gcc atc aca ggg ctc      240
Ser Ala Ser His Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu
              65                      70                      75                      80

cag gct gag gat gag gct gat tat tac tgc cag tcg tat gac acc agt      288
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Thr Ser
              85                      90                      95

cta cgt ggt ccg gtg ttc ggc gga ggg acc aag ctg acc gtc cta      333
Leu Arg Gly Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
              100                      105                      110

```

<210> 47  
 <211> 111  
 <212> PRT  
 <213> Human

<400> 47

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Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
                    5                      10                      15
Arg Val Thr Ile Ser Cys Ser Gly Ser Arg Ser Asn Ile Gly Ala His
                    20                      25                      30
Tyr Glu Val Gln Trp Tyr Gln Gln Phe Pro Gly Ala Ala Pro Lys Leu
                    35                      40                      45
Leu Ile Tyr Gly Asp Thr Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
                    50                      55                      60
Ser Ala Ser His Ser Gly Thr Ser Ala Ser Leu Ala Ile Thr Gly Leu
                    65                      70                      75                      80
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Thr Ser
                    85                      90                      95
Leu Arg Gly Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
                    100                      105                      110

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<210> 48  
 <211> 333  
 <212> DNA  
 <213> Human

<400> 48

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cag tct gtc gtg acg cag ccg ccc tca gtg tct ggg gcc cca ggg cag      48
Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
                    5                      10                      15
agg gtc acc atc tcc tgc act ggg agc agc tcc aac atc ggg aca ggt      96
Arg Val Thr Ile Ser Cys Thr Gly Ser Ser Ser Asn Ile Gly Thr Gly
                    20                      25                      30
tat gat gta cat tgg tac cag cag gtt cca gga tca gcc ccc aaa ctc      144
Tyr Asp Val His Trp Tyr Gln Gln Val Pro Gly Ser Ala Pro Lys Leu
                    35                      40                      45
ctc atc tat gct tac acc aat cgg ccc tca ggg gtc cct gac cga ttc      192
Leu Ile Tyr Ala Tyr Thr Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
                    50                      55                      60
tct ggc tcc aag tct ggc atg tca gcc tcc ctg gtc atc ggt ggt ctc      240
Ser Gly Ser Lys Ser Gly Met Ser Ala Ser Leu Val Ile Gly Gly Leu
                    65                      70                      75                      80
cag gct gag gat gag gct gat tat tac tgc cag tcc ttt gac gac agc      288
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Phe Asp Asp Ser
                    85                      90                      95
ctg aat ggt ctt gtc ttc gga cct ggg acc tcg gtc acc gtc ctc      333
Leu Asn Gly Leu Val Phe Gly Pro Gly Thr Ser Val Thr Val Leu
                    100                      105                      110

```

<210> 49  
 <211> 111  
 <212> PRT  
 <213> Human

<400> 49

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Gln Ser Val Val Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
                    5                      10                      15

Arg Val Thr Ile Ser Cys Thr Gly Ser Ser Ser Asn Ile Gly Thr Gly
                20                      25                      30

Tyr Asp Val His Trp Tyr Gln Gln Val Pro Gly Ser Ala Pro Lys Leu
                35                      40                      45

Leu Ile Tyr Ala Tyr Thr Asn Arg Pro Ser Gly Val Pro Asp Arg Phe
                50                      55                      60

Ser Gly Ser Lys Ser Gly Met Ser Ala Ser Leu Val Ile Gly Gly Leu
                65                      70                      75                      80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Phe Asp Asp Ser
                85                      90                      95

Leu Asn Gly Leu Val Phe Gly Pro Gly Thr Ser Val Thr Val Leu
                100                      105                      110

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<210> 50  
 <211> 333  
 <212> DNA  
 <213> Human

<400> 50

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cag tct gtg ttg acg cag ccg ccc tca gtg tct ggg gcc cca ggg cag      48
Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
                    5                      10                      15

agg gtc acc atc tcc tgc act ggg agc cac tcc aac ttc ggg gca ggt      96
Arg Val Thr Ile Ser Cys Thr Gly Ser His Ser Asn Phe Gly Ala Gly
                20                      25                      30

act gat gtc cat tgg tac caa cac ctt cca gga aca gcc ccc aga ctc      144
Thr Asp Val His Trp Tyr Gln His Leu Pro Gly Thr Ala Pro Arg Leu
                35                      40                      45

ctc att cat gga gac act cat cgg ccc tcc ggg gtc gct gac cga ttc      192
Leu Ile His Gly Asp Thr His Arg Pro Ser Gly Val Ala Asp Arg Phe
                50                      55                      60

tct ggc tcc agg tct ggc gcc tca gcc tcc ctg gcc atc act ggg ctc      240
Ser Gly Ser Arg Ser Gly Ala Ser Ala Ser Leu Ala Ile Thr Gly Leu
                65                      70                      75                      80

cgg gtt gag gat gag gct gat tat tac tgt cag tcg tat gac tat ggc      288
Arg Val Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Gly
                85                      90                      95

ctg aga ggt tgg gtg ttc ggc ggc ggg acc aag ctg acc gtc ctt      333
Leu Arg Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
                100                      105                      110

```

<210> 51  
 <211> 111  
 <212> PRT  
 <213> Human

<400> 51

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Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Gly Ala Pro Gly Gln
      5              10              15
Arg Val Thr Ile Ser Cys Thr Gly Ser His Ser Asn Phe Gly Ala Gly
      20              25              30
Thr Asp Val His Trp Tyr Gln His Leu Pro Gly Thr Ala Pro Arg Leu
      35              40              45
Leu Ile His Gly Asp Thr His Arg Pro Ser Gly Val Ala Asp Arg Phe
      50              55              60
Ser Gly Ser Arg Ser Gly Ala Ser Ala Ser Leu Ala Ile Thr Gly Leu
      65              70              75              80
Arg Val Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Gly
      85              90              95
Leu Arg Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
      100              105              110

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<210> 52  
 <211> 321  
 <212> DNA  
 <213> Human

<400> 52

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gac atc cag atg acc cag tct cca tct tcc gtg tct gca tct ata gga      48
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Ile Gly
      5              10              15

gac aga gtc acc atc act tgt cgg gcg agt cag ggt att gac aac tgg      96
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Asp Asn Trp
      20              25              30

tta ggc tgg tat cag cag aaa cct ggg aaa gcc cct aaa ctc ctg atc      144
Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
      35              40              45

tac gat gca tcc aat ttg gac aca ggg gtc cca tca agg ttc agt gga      192
Tyr Asp Ala Ser Asn Leu Asp Thr Gly Val Pro Ser Arg Phe Ser Gly
      50              55              60

agt gga tct ggg aca tat ttt act ctc acc atc agt agc ctg caa gct      240
Ser Gly Ser Gly Thr Tyr Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala
      65              70              75              80

gaa gat ttt gca gtt tat ttc tgt caa cag gct aaa gct ttt cct ccc      288
Glu Asp Phe Ala Val Tyr Phe Cys Gln Gln Ala Lys Ala Phe Pro Pro
      85              90              95

act ttc ggc gga ggg acc aag gtg gac atc aaa      321
Thr Phe Gly Gly Gly Thr Lys Val Asp Ile Lys
      100              105

```

<210> 53  
 <211> 107  
 <212> PRT  
 <213> Human

<400> 53

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Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Ile Gly
              5              10              15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Asp Asn Trp
              20              25              30

Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
              35              40              45

Tyr Asp Ala Ser Asn Leu Asp Thr Gly Val Pro Ser Arg Phe Ser Gly
              50              55              60

Ser Gly Ser Gly Thr Tyr Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala
              65              70              75              80

Glu Asp Phe Ala Val Tyr Phe Cys Gln Gln Ala Lys Ala Phe Pro Pro
              85              90              95

Thr Phe Gly Gly Gly Thr Lys Val Asp Ile Lys
              100              105

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<210> 54  
 <211> 13  
 <212> PRT  
 <213> Human

<400> 54

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Thr Gly Ser His Ser Asn Phe Gly Ala Gly Thr Asp Val
              5              10

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<210> 55  
 <211> 7  
 <212> PRT  
 <213> Human

<400> 55

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Gly Asp Ser Asn Arg Pro Ser
              5

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<210> 56  
 <211> 11  
 <212> PRT  
 <213> Human

<400> 56

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Gln Ser Tyr Asp Tyr Gly Leu Arg Gly Trp Val
              5              10

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<210> 57

<211> 11  
<212> PRT  
<213> Human

<400> 57

Arg Ala Ser Gln Asn Ile Asn Asn Tyr Leu Asn  
5 10

<210> 58  
<211> 7  
<212> PRT  
<213> Human

<400> 58

Ala Ala Ser Thr Leu Gln Ser  
5

<210> 59  
<211> 9  
<212> PRT  
<213> Human

<400> 59

Gln Gln Tyr Ser Arg Tyr Pro Pro Thr  
5

<210> 60  
<211> 14  
<212> PRT  
<213> Human

<400> 60

Thr Gly Ser Ser Thr Asp Val Gly Asn Tyr Asn Tyr Ile Ser  
5 10

<210> 61  
<211> 7  
<212> PRT  
<213> Human

<400> 61

Asp Val Thr Ser Arg Pro Ser  
5

<210> 62  
<211> 10  
<212> PRT  
<213> Human

<400> 62

Asn Ser Tyr Ser Ala Thr Asp Thr Leu Val  
5 10

<210> 63  
<211> 14  
<212> PRT  
<213> Human

<400> 63

Thr Gly Gln Ser Ser Asn Ile Gly Ala Asp Tyr Asp Val His  
5 10

<210> 64  
<211> 7  
<212> PRT  
<213> Human

<400> 64

Gly His Asn Asn Arg Pro Ser  
5

<210> 65  
<211> 11  
<212> PRT  
<213> Human

<400> 65

Gln Ser Tyr Asp Ser Ser Leu Ser Gly Leu Val  
5 10

<210> 66  
<211> 10  
<212> PRT  
<213> Human

<400> 66

Arg Ala Ser Gln Asp Ile Ser Trp Leu Ala  
5 10

<210> 67  
<211> 7  
<212> PRT  
<213> Human

<400> 67

Ala Ala Ser Leu Leu Gln Ser  
5

<210> 68  
<211> 9  
<212> PRT  
<213> Human

<400> 68

Gln Gln Ala Asp Ser Phe Pro Pro Thr  
5

<210> 69  
<211> 11  
<212> PRT  
<213> Human

<400> 69

Arg Ala Ser Gln Ser Ile Lys Arg Trp Leu Ala  
                    5                    10

<210> 70  
<211> 7  
<212> PRT  
<213> Human

<400> 70

Ala Ala Ser Thr Leu Gln Ser  
                    5

<210> 71  
<211> 9  
<212> PRT  
<213> Human

<400> 71

Gln Gln Ala Asn Ser Phe Pro Pro Thr  
                    5

<210> 72  
<211> 14  
<212> PRT  
<213> Human

<400> 72

Ser Gly Ser Arg Ser Asn Ile Gly Ala His Tyr Glu Val Gln  
                    5                    10

<210> 73  
<211> 7  
<212> PRT  
<213> Human

<400> 73

Gly Asp Thr Asn Arg Pro Ser  
                    5

<210> 74  
<211> 11  
<212> PRT  
<213> Human

<400> 74

Gln Ser Tyr Asp Thr Ser Leu Arg Gly Pro Val

5

10

<210> 75  
<211> 14  
<212> PRT  
<213> Human

&lt;400&gt; 75

Thr Gly Ser Ser Ser Asn Ile Gly Thr Gly Tyr Asp Val His  
5 10

<210> 76  
<211> 7  
<212> PRT  
<213> Human

&lt;400&gt; 76

Ala Tyr Thr Asn Arg Pro Ser  
5

<210> 77  
<211> 11  
<212> PRT  
<213> Human

&lt;400&gt; 77

Gln Ser Phe Asp Asp Ser Leu Asn Gly Leu Val  
5 10

<210> 78  
<211> 14  
<212> PRT  
<213> Human.

&lt;400&gt; 78

Thr Gly Ser His Ser Asn Phe Gly Ala Gly Thr Asp Val His  
5 10

<210> 79  
<211> 7  
<212> PRT  
<213> Human

&lt;400&gt; 79

Gly Asp Thr His Arg Pro Ser  
5

<210> 80  
<211> 11  
<212> PRT  
<213> Human

&lt;400&gt; 80

Gln Ser Tyr Asp Tyr Gly Leu Arg Gly Trp Val  
                                   5                                  10

<210> 81  
 <211> 11  
 <212> PRT  
 <213> Human

<400> 81

Arg Ala Ser Gln Gly Ile Asp Asn Trp Leu Gly  
                                   5                                  10

<210> 82  
 <211> 7  
 <212> PRT  
 <213> Human

<400> 82

Asp Ala Ser Asn Leu Asp Thr  
                                   5

<210> 83  
 <211> 9  
 <212> PRT  
 <213> Human

<400> 83

Gln Gln Ala Lys Ala Phe Pro Pro Thr  
                                   5

<210> 84  
 <211> 2351  
 <212> DNA  
 <213> Human

<400> 84

ggtagccgag aaagaaccgg ctcccgagtt ctggggcattt cgcccggctc gaggtgcagg	59
atg cag agc aag gtg ctg ctg gcc gtc gcc ctg tgg ctc tgc gtg gag	107
Met Gln Ser Lys Val Leu Leu Ala Val Ala Leu Trp Leu Cys Val Glu	
5                                  10                                  15	
acc cgg gcc gcc tct gtg ggt ttg cct agt gtt tct ctt gat ctg ccc	155
Thr Arg Ala Ala Ser Val Gly Leu Pro Ser Val Ser Leu Asp Leu Pro	
20                                  25                                  30	
agg ctc agc ata caa aaa gac ata ctt aca att aag gct aat aca act	203
Arg Leu Ser Ile Gln Lys Asp Ile Leu Thr Ile Lys Ala Asn Thr Thr	
35                                  40                                  45	
ctt caa att act tgc agg gga cag agg gac ttg gac tgg ctt tgg ccc	251
Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro	
50                                  55                                  60	
aat aat cag agt ggc agt gag caa agg gtg gag gtg act gag tgc agc	299
Asn Asn Gln Ser Gly Ser Glu Gln Arg Val Glu Val Thr Glu Cys Ser	

65	70	75	80	
gat ggc ctc ttc	tgt aag aca ctc	aca att cca aaa	gtg atc gga aat	347
Asp Gly Leu Phe	Cys Lys Thr Leu	Thr Ile Pro Lys	Val Ile Gly Asn	
	85	90	95	
gac act gga gcc	tac aag tgc ttc	tac cgg gaa act	gac ttg gcc tcg	395
Asp Thr Gly Ala	Tyr Lys Cys Phe	Tyr Arg Glu Thr	Asp Leu Ala Ser	
	100	105	110	
gtc att tat gtc	tat gtt caa gat	tac aga tct cca	ttt att gct tct	443
Val Ile Tyr Val	Tyr Val Gln Asp	Tyr Arg Ser Pro	Phe Ile Ala Ser	
	115	120	125	
gtt agt gac caa	cat gga gtc gtg	tac att act gag	aac aaa aac aaa	491
Val Ser Asp Gln	His Gly Val Val	Tyr Ile Thr Glu	Asn Lys Asn Lys	
	130	135	140	
act gtg gtg att	cca tgt ctc ggg	tcc att tca aat	ctc aac gtg tca	539
Thr Val Val Ile	Pro Cys Leu Gly	Ser Ile Ser Asn	Leu Asn Val Ser	
	145	150	155	160
ctt tgt gca aga	tac cca gaa aag	aga ttt gtt cct	gat ggt aac aga	587
Leu Cys Ala Arg	Tyr Pro Glu Lys	Arg Phe Val Pro	Asp Gly Asn Arg	
	165	170	175	
att tcc tgg gac	agc aag aag ggc	ttt act att ccc	agc tac atg atc	635
Ile Ser Trp Asp	Ser Lys Lys Gly	Phe Thr Ile Pro	Ser Tyr Met Ile	
	180	185	190	
agc tat gct ggc	atg gtc ttc tgt	gaa gca aaa att	aat gat gaa agt	683
Ser Tyr Ala Gly	Met Val Phe Cys	Glu Ala Lys Ile	Asn Asp Glu Ser	
	195	200	205	
tac cag tct att	atg tac ata gtt	gtc gtt gta ggg	tat agg att tat	731
Tyr Gln Ser Ile	Met Tyr Ile Val	Val Val Val Gly	Tyr Arg Ile Tyr	
	210	215	220	
gat gtg gtt ctg	agt ccg tct cat	gga att gaa cta	tct gtt gga gaa	779
Asp Val Val Leu	Ser Pro Ser His	Gly Ile Glu Leu	Ser Val Gly Glu	
	225	230	235	240
aag ctt gtc tta	aat tgt aca gca	aga act gaa cta	aat gtg ggg att	827
Lys Leu Val Leu	Asn Cys Thr Ala	Arg Thr Glu Leu	Asn Val Gly Ile	
	245	250	255	
gac ttc aac tgg	gaa tac cct tct	tcg aag cat cag	cat aag aaa ctt	875
Asp Phe Asn Trp	Glu Tyr Pro Ser	Ser Lys His Gln	His Lys Lys Leu	
	260	265	270	
gta aac cga gac	cta aaa acc cag	tct ggg agt gag	atg aag aaa ttt	923
Val Asn Arg Asp	Leu Lys Thr Gln	Ser Gly Ser Glu	Met Lys Lys Phe	
	275	280	285	
ttg agc acc tta	act ata gat ggt	gta acc cgg agt	gac caa gga ttg	971
Leu Ser Thr Leu	Thr Ile Asp Gly	Val Thr Arg Ser	Asp Gln Gly Leu	
	290	295	300	
tac acc tgt gca	gca tcc agt ggg	ctg atg acc aag	aag aac agc aca	1019
Tyr Thr Cys Ala	Ala Ser Ser Gly	Leu Met Thr Lys	Lys Asn Ser Thr	
	305	310	315	320
ttt gtc agg gtc	cat gaa aaa cct	ttt gtt gct ttt	gga agt ggc atg	1067
Phe Val Arg Val	His Glu Lys Pro	Phe Val Ala Phe	Gly Ser Gly Met	

325										330					335					
gaa tct ctg gtg gaa gcc acg gtg ggg gag cgt gtc aga atc cct gcg	1115																			
Glu Ser Leu Val Glu Ala Thr Val Gly Glu Arg Val Arg Ile Pro Ala																				
340 345 350																				
aag tac ctt ggt tac cca ccc cca gaa ata aaa tgg tat aaa aat gga	1163																			
Lys Tyr Leu Gly Tyr Pro Pro Pro Glu Ile Lys Trp Tyr Lys Asn Gly																				
355 360 365																				
ata ccc ctt gag tcc aat cac aca att aaa gcg ggg cat gta ctg acg	1211																			
Ile Pro Leu Glu Ser Asn His Thr Ile Lys Ala Gly His Val Leu Thr																				
370 375 380																				
att atg gaa gtg agt gaa aga gac aca gga aat tac act gtc atc ctt	1259																			
Ile Met Glu Val Ser Glu Arg Asp Thr Gly Asn Tyr Thr Val Ile Leu																				
385 390 395 400																				
acc aat ccc att tca aag gag aag cag agc cat gtg gtc tct ctg gtt	1307																			
Thr Asn Pro Ile Ser Lys Glu Lys Gln Ser His Val Val Ser Leu Val																				
405 410 415																				
gtg tat gtc cca ccc cag att ggt gag aaa tct cta atc tct cct gtg	1355																			
Val Tyr Val Pro Pro Gln Ile Gly Glu Lys Ser Leu Ile Ser Pro Val																				
420 425 430																				
gat tcc tac cag tac ggc acc act caa acg ctg aca tgt acg gtc tat	1403																			
Asp Ser Tyr Gln Tyr Gly Thr Thr Gln Thr Leu Thr Cys Thr Val Tyr																				
435 440 445																				
gcc att cct ccc ccg cat cac atc cac tgg tat tgg cag ttg gag gaa	1451																			
Ala Ile Pro Pro Pro His His Ile His Trp Tyr Trp Gln Leu Glu Glu																				
450 455 460																				
gag tgc gcc aac gag ccc agc cat gct gtc tca gtg aca aac cca tac	1499																			
Glu Cys Ala Asn Glu Pro Ser His Ala Val Ser Val Thr Asn Pro Tyr																				
465 470 475 480																				
cct tgt gaa gaa tgg aga agt gtg gag gac ttc cag gga gga aat aaa	1547																			
Pro Cys Glu Glu Trp Arg Ser Val Glu Asp Phe Gln Gly Gly Asn Lys																				
485 490 495																				
att gaa gtt aat aaa aat caa ttt gct cta att gaa gga aaa aac aaa	1595																			
Ile Glu Val Asn Lys Asn Gln Phe Ala Leu Ile Glu Gly Lys Asn Lys																				
500 505 510																				
act gta agt acc ctt gtt atc caa gcg gca aat gtg tca gct ttg tac	1643																			
Thr Val Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr																				
515 520 525																				
aaa tgt gaa gcg gtc aac aaa gtc ggg aga gga gag agg gtg atc tcc	1691																			
Lys Cys Glu Ala Val Asn Lys Val Gly Arg Gly Glu Arg Val Ile Ser																				
530 535 540																				
ttc cac gtg acc agg ggt cct gaa att act ttg caa cct gac atg cag	1739																			
Phe His Val Thr Arg Gly Pro Glu Ile Thr Leu Gln Pro Asp Met Gln																				
545 550 555 560																				
ccc act gag cag gag agc gtg tct ttg tgg tgc act gca gac aga tct	1787																			
Pro Thr Glu Gln Glu Ser Val Ser Leu Trp Cys Thr Ala Asp Arg Ser																				
565 570 575																				
acg ttt gag aac ctc aca tgg tac aag ctt ggc cca cag cct ctg cca	1835																			
Thr Phe Glu Asn Leu Thr Trp Tyr Lys Leu Gly Pro Gln Pro Leu Pro																				

580										585					590					
atc	cat	gtg	gga	gag	ttg	ccc	aca	cct	gtt	tgc	aag	aac	ttg	gat	act	1883				
Ile	His	Val	Gly	Glu	Leu	Pro	Thr	Pro	Val	Cys	Lys	Asn	Leu	Asp	Thr					
595					600					605										
ctt	tgg	aaa	ttg	aat	gcc	acc	atg	ttc	tct	aat	agc	aca	aat	gac	att	1931				
Leu	Trp	Lys	Leu	Asn	Ala	Thr	Met	Phe	Ser	Asn	Ser	Thr	Asn	Asp	Ile					
610					615					620										
ttg	atc	atg	gag	ctt	aag	aat	gca	tcc	ttg	cag	gac	caa	gga	gac	tat	1979				
Leu	Ile	Met	Glu	Leu	Lys	Asn	Ala	Ser	Leu	Gln	Asp	Gln	Gly	Asp	Tyr					
625					630					635					640					
gtc	tgc	ctt	gct	caa	gac	agg	aag	acc	aag	aaa	aga	cat	tgc	gtg	gtc	2027				
Val	Cys	Leu	Ala	Gln	Asp	Arg	Lys	Thr	Lys	Lys	Arg	His	Cys	Val	Val					
645					650					655										
agg	cag	ctc	aca	gtc	cta	gag	cgt	gtg	gca	ccc	acg	atc	aca	gga	aac	2075				
Arg	Gln	Leu	Thr	Val	Leu	Glu	Arg	Val	Ala	Pro	Thr	Ile	Thr	Gly	Asn					
660					665					670										
ctg	gaa	aat	cag	acg	aca	agt	att	ggg	gaa	agc	atc	gaa	gtc	tca	tgc	2123				
Leu	Glu	Asn	Gln	Thr	Thr	Ser	Ile	Gly	Glu	Ser	Ile	Glu	Val	Ser	Cys					
675					680					685										
acg	gca	tct	ggg	aat	ccc	cct	cca	cag	atc	atg	tgg	tat	aaa	gat	aat	2171				
Thr	Ala	Ser	Gly	Asn	Pro	Pro	Pro	Gln	Ile	Met	Trp	Phe	Lys	Asp	Asn					
690					695					700										
gag	acc	ctt	gta	gaa	gac	tca	ggc	att	gta	ttg	aag	gat	ggg	aac	cgg	2219				
Glu	Thr	Leu	Val	Glu	Asp	Ser	Gly	Ile	Val	Leu	Lys	Asp	Gly	Asn	Arg					
705					710					715					720					
aac	ctc	act	atc	cgc	aga	gtg	agg	aag	gag	gac	gaa	ggc	ctc	tac	acc	2267				
Asn	Leu	Thr	Ile	Arg	Arg	Val	Arg	Lys	Glu	Asp	Glu	Gly	Leu	Tyr	Thr					
725					730					735										
tgc	cag	gca	tgc	agt	gtt	ctt	ggc	tgt	gca	aaa	gtg	gag	gca	ttt	ttc	2315				
Cys	Gln	Ala	Cys	Ser	Val	Leu	Gly	Cys	Ala	Lys	Val	Glu	Ala	Phe	Phe					
740					745					750										
ata	ata	gaa	ggg	gcc	cag	gaa	aag	acg	aac	ttg	gaa					2351				
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755					760															

&lt;210&gt; 85

&lt;211&gt; 764

&lt;212&gt; PRT

&lt;213&gt; Human

&lt;400&gt; 85

Met	Gln	Ser	Lys	Val	Leu	Leu	Ala	Val	Ala	Leu	Trp	Leu	Cys	Val	Glu
5					10					15					

Thr	Arg	Ala	Ala	Ser	Val	Gly	Leu	Pro	Ser	Val	Ser	Leu	Asp	Leu	Pro
20					25					30					

Arg	Leu	Ser	Ile	Gln	Lys	Asp	Ile	Leu	Thr	Ile	Lys	Ala	Asn	Thr	Thr
35					40					45					

Leu	Gln	Ile	Thr	Cys	Arg	Gly	Gln	Arg	Asp	Leu	Asp	Trp	Leu	Trp	Pro
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

50					55					60					
Asn	Asn	Gln	Ser	Gly	Ser	Glu	Gln	Arg	Val	Glu	Val	Thr	Glu	Cys	Ser
65					70					75					80
Asp	Gly	Leu	Phe	Cys	Lys	Thr	Leu	Thr	Ile	Pro	Lys	Val	Ile	Gly	Asn
				85					90					95	
Asp	Thr	Gly	Ala	Tyr	Lys	Cys	Phe	Tyr	Arg	Glu	Thr	Asp	Leu	Ala	Ser
			100					105					110		
Val	Ile	Tyr	Val	Tyr	Val	Gln	Asp	Tyr	Arg	Ser	Pro	Phe	Ile	Ala	Ser
		115					120					125			
Val	Ser	Asp	Gln	His	Gly	Val	Val	Tyr	Ile	Thr	Glu	Asn	Lys	Asn	Lys
		130				135					140				
Thr	Val	Val	Ile	Pro	Cys	Leu	Gly	Ser	Ile	Ser	Asn	Leu	Asn	Val	Ser
145					150					155					160
Leu	Cys	Ala	Arg	Tyr	Pro	Glu	Lys	Arg	Phe	Val	Pro	Asp	Gly	Asn	Arg
				165					170					175	
Ile	Ser	Trp	Asp	Ser	Lys	Lys	Gly	Phe	Thr	Ile	Pro	Ser	Tyr	Met	Ile
			180					185					190		
Ser	Tyr	Ala	Gly	Met	Val	Phe	Cys	Glu	Ala	Lys	Ile	Asn	Asp	Glu	Ser
		195					200					205			
Tyr	Gln	Ser	Ile	Met	Tyr	Ile	Val	Val	Val	Val	Gly	Tyr	Arg	Ile	Tyr
		210				215					220				
Asp	Val	Val	Leu	Ser	Pro	Ser	His	Gly	Ile	Glu	Leu	Ser	Val	Gly	Glu
225					230					235					240
Lys	Leu	Val	Leu	Asn	Cys	Thr	Ala	Arg	Thr	Glu	Leu	Asn	Val	Gly	Ile
				245					250					255	
Asp	Phe	Asn	Trp	Glu	Tyr	Pro	Ser	Ser	Lys	His	Gln	His	Lys	Lys	Leu
			260					265					270		
Val	Asn	Arg	Asp	Leu	Lys	Thr	Gln	Ser	Gly	Ser	Glu	Met	Lys	Lys	Phe
		275					280					285			
Leu	Ser	Thr	Leu	Thr	Ile	Asp	Gly	Val	Thr	Arg	Ser	Asp	Gln	Gly	Leu
		290				295					300				
Tyr	Thr	Cys	Ala	Ala	Ser	Ser	Gly	Leu	Met	Thr	Lys	Lys	Asn	Ser	Thr
305					310					315					320
Phe	Val	Arg	Val	His	Glu	Lys	Pro	Phe	Val	Ala	Phe	Gly	Ser	Gly	Met
				325					330					335	
Glu	Ser	Leu	Val	Glu	Ala	Thr	Val	Gly	Glu	Arg	Val	Arg	Ile	Pro	Ala
			340					345					350		
Lys	Tyr	Leu	Gly	Tyr	Pro	Pro	Pro	Glu	Ile	Lys	Trp	Tyr	Lys	Asn	Gly
		355					360					365			
Ile	Pro	Leu	Glu	Ser	Asn	His	Thr	Ile	Lys	Ala	Gly	His	Val	Leu	Thr
		370				375					380				
Ile	Met	Glu	Val	Ser	Glu	Arg	Asp	Thr	Gly	Asn	Tyr	Thr	Val	Ile	Leu
385					390					395					400

Thr Asn Pro Ile Ser Lys Glu Lys Gln Ser His Val Val Ser Leu Val  
 405 410 415  
 Val Tyr Val Pro Pro Gln Ile Gly Glu Lys Ser Leu Ile Ser Pro Val  
 420 425 430  
 Asp Ser Tyr Gln Tyr Gly Thr Thr Gln Thr Leu Thr Cys Thr Val Tyr  
 435 440 445  
 Ala Ile Pro Pro Pro His His Ile His Trp Tyr Trp Gln Leu Glu Glu  
 450 455 460  
 Glu Cys Ala Asn Glu Pro Ser His Ala Val Ser Val Thr Asn Pro Tyr  
 465 470 475 480  
 Pro Cys Glu Glu Trp Arg Ser Val Glu Asp Phe Gln Gly Gly Asn Lys  
 485 490 495  
 Ile Glu Val Asn Lys Asn Gln Phe Ala Leu Ile Glu Gly Lys Asn Lys  
 500 505 510  
 Thr Val Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr  
 515 520 525  
 Lys Cys Glu Ala Val Asn Lys Val Gly Arg Gly Glu Arg Val Ile Ser  
 530 535 540  
 Phe His Val Thr Arg Gly Pro Glu Ile Thr Leu Gln Pro Asp Met Gln  
 545 550 555 560  
 Pro Thr Glu Gln Glu Ser Val Ser Leu Trp Cys Thr Ala Asp Arg Ser  
 565 570 575  
 Thr Phe Glu Asn Leu Thr Trp Tyr Lys Leu Gly Pro Gln Pro Leu Pro  
 580 585 590  
 Ile His Val Gly Glu Leu Pro Thr Pro Val Cys Lys Asn Leu Asp Thr  
 595 600 605  
 Leu Trp Lys Leu Asn Ala Thr Met Phe Ser Asn Ser Thr Asn Asp Ile  
 610 615 620  
 Leu Ile Met Glu Leu Lys Asn Ala Ser Leu Gln Asp Gln Gly Asp Tyr  
 625 630 635 640  
 Val Cys Leu Ala Gln Asp Arg Lys Thr Lys Lys Arg His Cys Val Val  
 645 650 655  
 Arg Gln Leu Thr Val Leu Glu Arg Val Ala Pro Thr Ile Thr Gly Asn  
 660 665 670  
 Leu Glu Asn Gln Thr Thr Ser Ile Gly Glu Ser Ile Glu Val Ser Cys  
 675 680 685  
 Thr Ala Ser Gly Asn Pro Pro Pro Gln Ile Met Trp Phe Lys Asp Asn  
 690 695 700  
 Glu Thr Leu Val Glu Asp Ser Gly Ile Val Leu Lys Asp Gly Asn Arg  
 705 710 715 720  
 Asn Leu Thr Ile Arg Arg Val Arg Lys Glu Asp Glu Gly Leu Tyr Thr  
 725 730 735  
 Cys Gln Ala Cys Ser Val Leu Gly Cys Ala Lys Val Glu Ala Phe Phe

			740					745				750
Ile	Ile	Glu	Gly	Ala	Gln	Glu	Lys	Thr	Asn	Leu	Glu	
		755					760					

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/US02/06762

**A. CLASSIFICATION OF SUBJECT MATTER**

IPC(7) : A61K 39/40, 39/42, 39/395, 51/00

US CL : 424/1.11, 135.1, 142.1, 143.1, 155.1, 174.1

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 424/1.11, 135.1, 142.1, 143.1, 155.1, 174.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)  
WEST, Medline, Biosis**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US 5,840,301 A (ROCKWELL et al.) 24 November 1999 (24.11.1999), see entire document.	1-4 and 33-37
Y	US 5,861,499 A (ROCKWELL et al.) 19 January 1999 (19.01.99), see entire document.	1-4 and 33-37
Y,E	US 6,365,157 B2 (ROCKWELL et al.) 02 April 2002 (02.04.02), see entire document.	1-4 and 33-37
Y,P	US 2002/0019414 A1 (ALTMANN et al.) 14 February 2002 (14.02.02), see entire document.	1-4 and 33-37
Y,P	US 6,297,238 B1 (DOYLE et al.) 02 October 2001 (02.10.01), see entire document.	1-4 and 33-37
Y	BENNINGTON. Saunders Dictionary & Encyclopedia of Laboratory Medicine and Technology (Philadelphia, PA, W. B. Saunders Company, 1984), pages 289-291, 1288, and 1289.	1-4 and 33-37
Y	BUCHDUNGER et al. 4,5-Dianilinophthalimide: a protein-tyrosine kinase inhibitor with selectivity for the epidermal growth factor receptor signal transduction pathway and potent in vivo antitumor activity. Proceedings of the National Academy of Sciences of the United States of America. 15 March 1994, Vol. 91, No. 6, pages 2334-8.	1-4 and 33-37

☐ Further documents are listed in the continuation of Box C.☐ See patent family annex.

## \* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&amp;" document member of the same patent family

Date of the actual completion of the international search

29 May 2002 (29.05.2002)

Date of mailing of the international search report

02 JUL 2002

Name and mailing address of the ISA/US

Commissioner of Patents and Trademarks

Box PCT

Washington, D.C. 20231

Facsimile No. (703)305-3230

Authorized officer

Jeffrey Stucker

Telephone No. 703-308-0196

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US02/06762

## Box I Observations where certain claims were found unsearchable (Continuation of Item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claim Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☐ Claim Nos.:  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☒ Claim Nos.: 5-32,38 and 39  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of Item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

☐  
☐

The additional search fees were accompanied by the applicant's protest.

No protest accompanied the payment of additional search fees.